

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(4517)

<223> v=yes-1 yamaguchi sarcoma viral oncogene
homolog 1 (YES1) gene.

<400> 304

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198/292

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gagtctcgct ctgccgcca ggctggagtg cagtggcgcg atctcgctc actgcaagct 3240
ctgectccca ggtcacacc attctcctgc ctgagcctcc cgagtagctg ggactacagg 3300
tgccaccac cagcctggc tagttttttg tatttttagt agagacgcag tttcacctg 3360
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<210> 305

<211> 459

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(459)

<223> 3' terminal sequence. interferon -induced protein 75, 52kd (IFI75) gene.

<400> 305

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ctttatctct tatttggggg atcaggttgt cactggccac ttgcacagt ctagtgagga 180
ggctgggcat ctctctctgag tcttctttcg cattcatttt ggangttaac ttgtcatttg 240
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gggctgggag actcactcag gatctcatcg ctttctggg agg atgttc agggctcact 360
gactcttggg cgcacaaggt gaaacagctt ggtttgaagg gggttnttgg tngggggcaa 420
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<210> 306

<211> 370

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature
 <222> (1)..(370)
 <223> 5' terminal sequence. interferon -induced
 protein 75, 52kd (IFI75) gene.

<400> 306
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 aatcttgagg agtgatccct gtccc agccc ctggaaagg anggaaacga caaactcaaa 120
 gtccaggatg ttcacatga caagagccat ggaagaggct ctttttcagc acttcatgca 180
 ccagaagctg gggatcgct atgccatata caagccattt cccttctttg aaggcctcct 240
 agacaactcc atcatcacta agagaatgta catggaatct ctggaagcct gtagaaattt 300
 gatccctgta tccagagtgg tgcacaacat tctcacccaa ctgggagagg actttttaac 360
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<210> 307
 <211> 1541
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(1541)
 <223> interferon -induced protein 75, 52kd (IFI75)
 gene.

<400> 307
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 tccaggaagg aagaagcact tcagtgaacca atgacaagtt aacatccaaa atgaatgcgg 180
 aagaagactc agaagagatg cccagcctcc tctactagcac tgtgcaagtg gccagtgaca 240
 acctgatccc ccaaataaga gataaagaag acctcaaga gatgcccac tctcccttgg 300
 gctctatgcc agagataaga gataattctc cagaacccaa t gaccagaa gagccccagg 360
 aggtgtccag cacacettca gacaagaaag gaaagaaaag aaaaagatgt atctggtcaa 420
 ctccaaaaag gagacataag aaaaaaagcc tcccagagg gacagcctca tctagacacg 480
 gaatccaaaa gaagctcaaa aggggtggatc aggttcctca aaagaaagat gactcaactt 540
 gtaactccac ggtagagaca agggcccaaa aggcgagaaac tgaatgtgcc cgaaagtcca 600
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<210> 308
 <211> 416
 <212> DNA

200/292

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(416)

<223> 3' terminal sequence. v-myb avian
myeloblastosis viral oncogene homolog -like 2
(MYBL2) gene.

<400> 308

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agggcaggnc tgacgtgggc ttgggagaag ctgacggagc tccctgtggc cttggggagg 180
gaaccaggca gacctgggaa gtggaacttt gttgttagca ccaggagccg cccacagctg 240
ggcttcggca acagggcagc acatggccct gttccttcca cctgagagt c tggggagggg 300
ctggtggcag aaggctccct gcaggaggtt caoctgaatg actctcagat tcacagaccc 360
cctnttgccc ccacaacccc tgtaaacatg agaatggggc tcgtgacacc ctnaac 416
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<210> 309

<211> 426

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(426)

<223> 5' terminal sequence. v-myb avian
myeloblastosis viral oncogene homolog -like 2
(MYBL2) gene.

<400> 309

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agctgatgat gtccacactg cccaagtctc tctccttgcc gacaactgcc cttcaaact 180
cttcagcct caccctgtca ggtatcaaag aagacaacag cttgctcaac cagggtttot 240
tgcaggccaa gcccgagaag gcagcagtgg ccagaagcc ccgaagc cac ttcacgacac 300
ctgcccctat gtccagtgcc tgggaagacg gtggcctgcg gggggaccag gggaccagct 360
tttcatggca ggagaaagcc cggcagcttc tggggccgct tgaagcccag ccacactttt 420
cgggac 426
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<210> 310

<211> 2627

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2627)

<223> v-myb avian myeloblastosis viral oncogene
homolog-like 2 (MYBL2) gene.

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<400> 310
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gcctcatctc agaccctgct taggatggg gatgtggcca ggggtgctcc tgtgctcacc 2580
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<210> 311

<211> 442

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(442)

<223> 3' terminal sequence. transforming growth factor, beta receptor iii (betaglycan, 300kd)

202/292

(TGFB3) gene.

<400> 311

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cccagactca aggagttggt aaagggttaa tagccagata gtagaaccag tgaggagatg 60
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taagcgatcc tctgtctaa ttgacacctt tgtctggatg cacacttctg accttgctgc 180
cacaacctgt ggggttctga tgtgtccctt gatgggtgct gccctcaggg actgcacct 240
gacaagtgtt aaggcaacat tcttttctg tgcccggggc caaaaccaat gctgatgacc 300
ttatcagctt cctgtttctt cccatacttg catacaccac tggcaaatg tcttaatggc 360
aaattttgta ttcttacag ggnctacag aaatttgaaa atgg accaaa ttcagggaac 420
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442

<210> 312

<211> 315

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(315)

<223> 5' terminal sequence. transforming growth factor, beta receptor iii (betaglycan, 300kd) (TGFB3) gene.

<400> 312

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taacaaggag gtatcactga gcttatttta gctgcaaagt ggcatcatat tattccattt 60
aatgaaattc acctcaagcc ctttttgaca tattaaatat atgggatata tttaa ggcaa 120
gagaagtaag gcaatccaaa tgagtgcctt ttccaatct cagcactgtc ttgngnga 180
tggtgacact attcagataa ccaactggag accgacagat ttgccatgca ttgcatctt 240
gctagagttt ggtttttatg aaagggccta ttttttttta agttgacata ttttgagtgg 300
gaaacactca ccta

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315

<210> 313

<211> 4208

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(4208)

<223> transforming growth factor, beta receptor iii (betaglycan, 300kd) (TGFB3) gene.

<400> 313

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gaaagaaccg catgagcctg acggcgcatg gtcttaacat caggctgtgc aggaagaagc 120
tatctgcaga tggatgccag cacacacaag gaagcagagc tctggcaaca ttgagtcaaa 180
gcaaggacac aacatcagag ggacggcaga gaatccttgt gtgtagtctt tgggtggcagt 240
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agcttcactg ttttgtcagg ctgtgccagc agaggcaca ctgggctgcc acaggagggtg 540

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203/292

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catgtcctga atctcgcact ggcgcagggg cctggccagc tacagagaga ggtcacactt 600
cacctgaate ccattctctc a gtccacatc caccacaagt ctgttgtgtt cctgctcaac 660
tccccacacc ccttgggtgtg gcattctgaag acagagagac ttgccactgg ggtctccaga 720
ctgttttttg tgtctgaggg ttctgtgtgc cagttttcat cagcaaactt ctccttgaca 780
gcagaaacag aagaaaggaa cttcccccat ggaaatgaac atctgttaaa ttgggccccg 840
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aatctcatta aatggaataa tatgatgcca ctttgcagct aaaataagct cagtataacc 4200

```

tccttggt

4208

<210> 314

<211> 468

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(468)

<223> 3' terminal sequence. peroxiredoxin 2
(PRDX2) gene.

<400> 314

```
tnnttttttt tttncacott tccctaatac ttnatnggtn acctctaggc ctgtgtgcgg 60
ctgggtgggc ttgggggagg gcgtcactat tcagcttcta ggtggaggca tgagaaggcc 120
ttggctaggc cctccagggt cccatactgt ggagtttga ggggcaggtc tggcctttcc 180
tgggtcagca tagggcacc aggtgggggn acaggtggac acccagcaca ggcacctagg 240
caggggcaca agtcantat cnttagcca gctaattgt ntttgg agaa atattccttg 300
ctgtcatcca cgttgggttt aatcgtgtca ctgccaggtt tccagccagc gggacaaant 360
ttcccatgt tcgtttgtgt attggaagg cctgggacca gccgcagagt tnatccacg 420
gagngtccca aaggnaaatc attaaacagt gatattggcn aaggaaaa 468
```

<210> 315

<211> 394

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(394)

<223> 5' terminal sequence. peroxiredoxin 2
(PRDX2) gene.

<400> 315

```
acttcaaggc cacagcgggt gttgatggcg ctttcaaaga ggtga agctg tcggactaca 60
aagggaagta cgtggctctc tttttctacc ctctggactt cacttttgtg tgccccaccg 120
agatcatcgc gttcagcaac cgtgcagagg acttccgcaa gctgggctgt gaagtgtgtg 180
gcgtctcggg tggactctca gttcaccac ctggettga tcaacacccc ccgaaagag 240
ggaggcttgg gccccctgaa catc cccctg ctgtgtgacg tgaccagaag cttgtctgag 300
gattacggcg tgctgaaaac agatgagggc attgctaaca ggggcctctt tatcatcgat 360
gggcaagggt gttcctttcg ccagatcaat gtta 394
```

<210> 316

<211> 937

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

205/292

<221> misc_feature
<222> (1)..(937)
<223> peroxiredoxin 2 (PRDX2) gene.

<400> 316
cgcgggcccca gggctcactt ggcgctgaga acgcgggtgc agcgtgtgat cgtccgtgcg 60
tctagccttt gccacgcag ctttcagtca tggcctcgg taacgcg cgc atcggaagc 120
cagccctga cttcaaggcc acagcgggtg ttgatggcgc cttcaaagag gtgaagctgt 180
cggactacaa agggaagtac gtggtcctct tttctaccc tctggacttc acttttgtgt 240
gccccaccga gatcatcgcg ttcacaaccg tgaagaggac ttccgcaaag ctgggctgtg 300
aagtgtggg cgtctcgggtg gaactcagt tcacccacct ggcttggatc aacaccccc 360
ggaaagaggg aggcttgggc cccttgaaca tccccctgct tgctgacgtg accagacgct 420
tgtctgagga ttacggcgtg ctgaaaaacg atgagggcac tgcttacagg ggctcttta 480
tcatcgatgg caagggtgtc cttogccaga tcaactgttaa tgatttgct gtgggacgct 540
cgtggatga ggctctgcgg ctggtccagg ccttccagta cacagacgag catggggaag 600
tttgtccggc tgcttgggaag cctggacgtg acacgattaa gccgaacgtg gatgacagca 660
aggaatatTT ctccaaacac aattaggctg gctaacgat agtgagcttg tgccccctgcc 720
taggtgcctg tgctgggtgt ccacctgtgc cccacctgg gtgcc ctatg ctgaccagc 780
aaaggccaga cctgccccctc caaaatccac agtatgggac cctggagggc tagcaaggcc 840
ttctcatgcc tccacctaga agctgaatag tgacgcctc cccaagccc acccagccgc 900
acacaggcct agaggtaacc aataaagtat tagggcc 937

<210> 317
<211> 451
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(451)
<223> 5' terminal sequence. v-fos fbj murine
osteosarcoma viral oncogene homolog (FOS) gene.

<400> 317
gctagcacca tgagctgaag accgagccct ttgatgactt cctgttccca gcatcatcca 60
ggcccagtgg ctctgagaca gcccgctcgg tgccagacat ggacctatct gggctcttct 120
atgcagcaga ctgggagcct ctgcacagtg gctccctggg gatggggccc atggcacagn 180
agctggagcc cctgtgcact ccggtggtca cctgtactcc cagctgcact gctta cacgt 240
cttcttctgt cttcacctac cccgaggctg actncttccc cagctgtgca gctgcccacc 300
gcaaggcagc agcagcaatg agccttctc tgactcgttc agctnaccga cgggtgctggc 360
cctgtgaggg ggcaggggaa ggggaggcag nccgcaacna caagttgcca ttgtccgagt 420
tngttgattt anagagagga gaaacaaatt t 451

<210> 318
<211> 2084
<212> DNA/RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(2084)
<223> v-fos fbj murine osteosarcoma viral oncogene
homolog (FOS) gene.

<400> 318

```
aaccgcatct gcagcgagca actgagaagc caagactgag ccggcgggccg cggcgcgagcg 60
aacgagcagt gaccgtgctc ctaccagct ctgcttcaca gcgcccacct gtctccgccc 120
ctcgccccct cgcccggtt tgccaaaccg ccaagatgat gttctcgggc ttcaacgcag 180
actacgagggc gtcate ctcc cgtgcagca gcgcgtcccc ggccggggat agcctctctt 240
actaccactc acccgagac tctttctcca gcatgggctc gcctgtcaac gcgcaggact 300
tctgcacgga cctggccgtc tccagtgcc acttcattcc caggttcaat gccatctcga 360
ccagtccgga cctgcagtgg ctggtgcagc ccgcccctgt ctctctgtg gcccctgc 420
agaccagagc cctcaccct ttggagtcc ccgcccctc cgtgggggt tactccagg 480
ctggcggtgt gaagaccatg acaggaggcc gagcgagag cattggcagg aggggcaagg 540
tggaacagtt atctccagaa gaagaagaga aaaggagaat ccgaaggga aggaataaga 600
tggtgcagc caaatgccgc aaccggagga gggagc tgac tgatacactc caagcgaga 660
cagaccaact agaagatgag aagtctgctt tgacagaccg gattgccaac ctgctgaagg 720
agaaggaaaa actagagttc atcttgccag ctaccgacc tgctgcaag atccctgatg 780
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gtccctggg gatggggccc atggccacag agctggagcc cctgtgact ccggtggtca 1140
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ccctagaggg ttctgtaga cctagggagg accttatctg tgcgtgaaac acaccaggct 1440
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ctgattagaa atgaccaata ttatactaag aaaagatacg actttatttt ctggtagata 1800
gaaataaata gctatatcca tgtactgtag tttttcttca acatcaatgt tcattgtaat 1860
gttactgatc atgcattgtt gaggtgtct gaatgttctg acattaacag ttttccatga 1920
aaacgtttta ttgtgttt tt aatttattta ttaagatgga ttctcagata tttatatttt 1980
tattttattt ttttctacct tgaggtcttt tgacatgtgg aaagtgaatt tgaatgaaaa 2040
atttaagcat tgtttgctta ttgttccaag acattgtcaa taaa 2084
```

<210> 319

<211> 240

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(240)

<223> 3' terminal sequence. retinoblastoma -binding
protein 7 (RBBP7) gene.

<400> 319

```
ctgcaaagcc aatcaagaag tggtggaagg aaaaagtgtg aaagttattc ttgcatattt 60
gggaacagca agcacttagt ttgagaaaat gaggacttaa aacagttgan tcaaaggcaa 120
taccctgcta cttgtattta aaatcaatgg tgatgttatt tcttangcaa cattcttctc 180
ttccctaata gctacaatnt gatacagtag gcaacagctc acttgaaagt gctagantca 240
```

<210> 320

207/292

<211> 457
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(457)
 <223> 5' terminal sequence. retinoblastoma -binding
 protein 7 (RBBP7) gene.

<400> 320
 agatgtttga agatactgtg gaggagcgtg tcatcaatga agaatat aaa atctggaaga 60
 agaatacacc gtttctatat gacctgggta tgacctatgc tcttcagtgg cccagtctta 120
 ccgttcagtg gcttcctgaa gtgactaaac ctgaaggaaa agattatgcc cttcattggc 180
 tagtgctggg gactcatacg tctgatgagc agaatcatct ggtgggttgc cgagtacata 240
 ttcccaatga tgatgcacag tttgat gctt cccattgtga cagtgcacaag ggtgaatttg 300
 gtggcctttg ttctgtaaca ggnaaaattg aatgtgaaat taaaatcaat tcacgaagga 360
 gaagttaaac cgtgctcggt aacatggcgc cagantcctt cacatccatt gcttacaaan 420
 acaccctctt gcttgatggt gttggnnttt tgactat 457

<210> 321
 <211> 1946
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(1946)
 <223> retinoblastoma -binding protein 7 (RBBP7)
 gene.

<400> 321
 gcctcgtcag ctgcctgggc gggc tgggag ggcgcgggtg aaaagtctcg ttccaagttt 60
 ggagagagag agaagagcgc ctacagacctc ggtacccgcg agcggggagg aggcaggaaa 120
 gaaggacgcg gcgtctgggg agcaccacag cagcaagacg gggcccgggc tttcgacagt 180
 ggggagtgag acgcgcttgg gaaaggcagg agcgcacgcg gtcgggctgc tcttggttaa 240
 cgagaggagt ccgaggcggc ggcgaggggc gaacgacccg acgcaagatg gcgagtaaa 300
 agatgtttga agatactgtg gaggagcgtg tcatcaatga agaataataa atctggaaga 360
 agaatacacc gtttctatat gacctgggta tgacctatgc tcttcagtgg cccagtctta 420
 ccgttcagtg gcttcctgaa gtgactaaac ctgaaggaaa agatt atgcc cttcattggc 480
 tagtgctggg gactcatacg tctgatgagc agaatcatct ggtgggttgc cgagtacata 540
 ttcccaatga tgatgcacag tttgatgctt cccattgtga cagtgcacaag ggtgaatttg 600
 gtggcctttg ttctgtaaca ggaaaaattg aatgtgaaat taaaatcaat cacgaaggag 660
 aagtaaaccg tgctcgctac atg ccgcaga atcctcacat cattgctaca aaaacaccat 720
 cttctgatgt gttgggtttt gactatacaa aacaccctgc taaaccagac ccaagtggag 780
 aatgtaatcc tgatctcaga ttaagaggtc accagaagga aggctatggt ctctcctgga 840
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 atataaacgc aggacaaaaa gaaggcaaaa ttgtggatgc taaagccatc tttactggcc 960
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 gtcacttggg ggatgcgcac actgcccgaag tcaactgcct c tcattcaat cctacagcg 1140
 aatttattct agccaccggc tctgcggata agaccgtagc tttatgggat ctgcgtaact 1200
 taaaattaaa actccatacc ttogaatctc ataaagatga aattttccag gtccactggg 1260
 ctccacataa tgaaactatt ctggcttcaa gtggtactga ccgcgcctg aatgtgtggg 1320

208/292

```

attttaagtaa aattggggaa gaacaatcag cagaagatgc agaagatggg cctccagaac 1380
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agccttgggt catttgctca gtgtctgagg ataacatcat gcagatatgg caaatggctg 1500
aaaatattta caatgatgaa gagtcagatg tcacgacatc cgaactggag ggaca aggat 1560
cttaaaccga aagtacgaga aatgtttctg ttgaatgtaa tgctacatga atgcttgatt 1620
tatcaagcgc caaaaaggca ttgtatagta ggaaatgtaa gtgggggtggc ttatggcttc 1680
tttatcctct gattctagca ctttcaagtg agctgttgcg tactgtatca tattgtagct 1740
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agcaggggat tgcctttgat tcaactgttt taagtctcct ttttctcaaa ctaagtgcct 1860
gctgttccca aatatgcaag aataactttt acactttttc cttccaacac ttcttgattg 1920
gctttgcaga aataaagttt taaaat 1946

```

<210> 322

<211> 365

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(365)

<223> 3' terminal sequence. kiaa1075 protein
(KIAA1075) gene.

<400> 322

```

tatagaaatt ctttattatt agacaaaaat agactctctt ttttccccta ttcatgtgat 60
cctactctga atctctgctc agaggaggca gtgactcgct cccacccct ctcccatccc 120
tgccgtgctg gcacctgcag ctgggtggaa ctggcagggg ctgatccct gggagggctg 180
acgtttctct gcaggtgggg ctgcctgctc tcttcgggg ctcaactgct gatgcctcct 240
caccctcacc acaccatct ctgccatctg ctcacatggc aagggtcctc ttgtggggcc 300
tggtccactt taacttaggc agggctgggg ggcgggggaa gggagaggca gtgttcccag 360
gggcc 365

```

<210> 323

<211> 400

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(400)

<223> 5' terminal sequence. kiaa1075 protein
(KIAA1075) gene.

<400> 323

```

tcaagggctg cccagtgag ccctactttg gcagcctgtc cgccttgggc tcccagcact 60
ccatctcccc catctccctg ccctgctgcc tgcgattcc cagcaaagat cctctggaag 120
agaccccaga ggtccagtg cccaccaaca tgagcacagc gncagacctc ctgcgtcagg 180
gtgctgcctg caggtngctc tacttgacct cagtggagac agagtcaact acgggcccc 240
aagctgtggc cggggccagc tctgcagctc tgagctgta g cccccgccg acaccagctg 300
ttgtccactt caaggtgtca gccaggga ttnacactga cgggacaacc aaaggaagct 360
ctttnttttc gccgccatta tccagtggaa cagcatcacc 400

```

209/292

<210> 324
 <211> 489
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(489)
 <223> 5' terminal sequence. atp-binding cassette,
 sub-family c (cftr/mrp), member 5 (ABCC5) gene.

<400> 324
 nttcggcaca gnaagataca actctgtgct gaacagctgc tgccctgaggc tgacctggcc 60
 attcttccca gcagcgacct gacggagatt ggagagcgag tancaacctg agcgggtgggc 120
 agcgcagagg atcagccttg cccgggcctt gtatagtgc aggagcatct acatcctgga 180
 cgacccctc agtgccttag atgcccattg ggaanccaca tncctcaata gtgctatccg 240
 gaaacatctc aagtccaaga cagttctgtt tgttacccac cagt tacagt acctgggttg 300
 actgtgatga agtgatcttc atgaaagagg gctgtattta cgggaaagag ggcaccntg 360
 gaggaantg atggatttta aatggtgatt atggttacct ttttaattaa cntgttggtg 420
 ggggagagac accgccattg agntcatttc aaaaagggga accgtnggtt cacagaggag 480
 ttcacagtt 489

<210> 325
 <211> 5838
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(5838)
 <223> atp-binding cassette, sub-family c
 (cftr/mrp), member 5 (ABCC5) gene.

<400> 325
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<210> 326

<211> 385

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(385)

<223> 3' terminal sequence. cadherin 1, type 1,
e-cadherin (epithelial) (CDH1) gene.

<400> 326

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<210> 327

<211> 423

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(423)

<223> 5' terminal sequence. cadherin 1, type 1,
e-cadherin (epithelial) (CDH1) gene.

<400> 327

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<210> 328

<211> 4828

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(4828)

<223> cadherin 1, type 1, e-cadherin (epithelial)
(CDH1) gene.

<400> 328

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<210> 329

<211> 471

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(471)

<223> 5' terminal sequence. zinc finger protein
144 (mel-18) (ZNF144) gene.

<400> 329

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<210> 330

<211> 2227

<212> DNA/RNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2227)

<223> zinc finger protein 144 (mel -18) (ZNF144)
gene.

<400> 330

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aagcggctca ccttagccac ggtgcccacc cctccgagg gcaccaacac cagcggggcg 960
tccgagtgtg agtcagtcag cgacaaggct cccagccctg ccaccctgcc agccacctcc 1020
tcctccctgc ccagccagc caccctatcc catggctctc ccagttccca tgggcctcca 1080
gccaccacc ctacctccc cactccctc tcgacagcca gtggggccac c acagctgcc 1140
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atgtacatag gaaaaaccaa atatacatc ttattttcta tggaccaacc agattaattt 1440
aaatgccaca ggaaacaaac tttatgtgtg tgtgtatgtg tggaaaatgg tgttcatttt 1500
ttttgggggg ggtcttgtgt aatttgctgt ttttgggggt gcctggagat gaactggatg 1560
ggccactgga gtctcaataa agctctgcac catcctcgct gtttcccaag gcagggtggtg 1620
tgttgggggc ccttcagac ccaaagcttt aggcattgatt ccaactgget gcataatagga 1680
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aggccctgag aggt ctacgc tcttgagga gggctaaggc tttagcattg tgaagcgtg 2040
caccaccacc aaccttacc tcaccgggga accctacta gcaggactgg tgggtgagtc 2100
tcacctgggg cctagagtgg aagtgggggt gggttaacct cacacaagca cagatcccag 2160
actttgccag aggcacacag ggaattccgc cgatactgac gggctccagg agt cgtcgcc 2220
acactcg 2227

```

<210> 331
<211> 254
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(254)
<223> 3' terminal sequence. macrophage stimulating
1 (hepatocyte growth factor -like) (MST1) gene.

<400> 331
gcataaagag gaaacatggc tttatgtctg acaagaagtt ttgtcctccc caaggcatat 60
ggcatcaagg ctgggctaac ccagtctcat gaccttgtga atccagtcca caaacacaga 120
gacacgcgtg aagacagctg gccagcggga ccttgccat actcggttgg ggattataat 180
tccttccagg gacccagcag ttgtgggtaa agcaggcaag tgggcccccg tagtcaccct 240
cacaggcccc caca 254

<210> 332
<211> 362
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(362)
<223> 5' terminal sequence. macrophage stimulating
1 (hepatocyte growth factor -like) (MST1) gene.

<400> 332
gccatggncc tgggtgctaca cgatggaccc aaggacccca t tgcactact gtgccctgcg 60
acgctgcgct gatgaccagc cgccatcaat cctggacccc ccagaccagg tgcagtttga 120
gaagtgtggc aagagggtgg atcggtctga tcagcggcgt tccaagctgc gctggttggg 180
ggccatccgg gcaactcacc ctggacagtc agcttgccga atcggcaggg ccagcatttc 240
tgcggnggt ctctagtga ggagcagtn atactgactn cccggaagtg cttctcctcc 300
tnccatatnc ctctcacggg ctatgaggta tggttngggc ancctttttc cagaaccac 360
ag 362

<210> 333
<211> 2219
<212> DNA/RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(2219)
<223> macrophage stimulating 1 (hepatocyte growth
factor-like) (MST1) gene.

```

<400> 333
agccagaagg atgggggtggc tcccactcct gctgcttctg actcaatgct taggggtccc 6 0
tgggcagcgc tcgccattga atgacttcca agtgctccgg ggcacagagc tacagcacct 120
gctacatgcg gtgggtgccc ggcccttgga ggaggatgtg gcagatgctg aagagtgtgc 180
tggctcgtgt gggcccttaa tggactgccc ggcccttccac tacaacgtga gcagccatgg 240
ttgccaactg ctgccatgga ctcaacactc gccccacacg aggctgcggc gttctggggc 300
ctgtgacctc ttccagaaga aagactacgt acggacctgc atcatgaaca atggggttgg 360
gtaccggggc accatggcca cgaccgtggg tggcctgccc tgccaggctt ggagccacaa 420
gttcccgaat gatcacaagt acacgcccac tctccggaat ggcttgaag agaacttctg 480
ccgtaaccct gatggcgac c cggagggtcc ttggtgctac acaacagacc ctgctgtgcg 540
cttccagagc tgcggcatca aatcctgccc ggaggcgcgc tgtgtctggg gcaatggcga 600
ggaataccgc ggcgcggtag accgcacgga gtcagggcgc gagtgccagc gctgggatct 660
tcagcacccc caccagcacc ccttcgagcc gggaagtgc ctgcaccaag gtctggacga 720
caactattgc cggatcctg acggctccga gcggccatgg tgctacacta cggatccgca 780
gatcgagcga gaggttctgt acctcccccg ctgcgggtcc gaggcacagc cccgccaaga 840
ggccacaact gtcagctgct tccgcgggaa ggggtgagggc taccggggca cagccaatac 900
caccactgcg ggcgtacctt gccagcgttg ggacgcgc a atcccgcatc agcaccgatt 960
tacgccagaa aaatacgctg gcaaagacct tcgggagaac ttctgcgga accccgacgg 1020
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gatccggcgt tgtacagacg acgtgcggcc ccaggactgc taccacggcg caggggagca 1140
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gacgcgcgac aagccgcagt tcacgtttac ctccgaaccg catgcacaac tggaggagaa 1260
cttctgcgcg aaccagatg gggatagcca tgggcccctg tgctacacga tggaccaag 1320
gaccccatte gactactgtg ccttcgacg ctgcgctgat gaccagccgc cat caatcct 1380
ggacccccca gaccaggtgc agtttgagaa gtgtggcaag aggggtggatc ggctggatca 1440
gcggcggttc aagctgcgcg tgggtggggg ccacccgggc aactcaccct ggacagtacg 1500
cttgcggaat cggcagggcc agcatttctg cggggggtct ctagtgaagg agcagtggat 1560
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gttgggcacc ctgttccaga acccacagca tggagagcca agcctacagc ggggtcccagt 1680
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tccagggacc aagtgtgaga ttgcaggctg ggggtgagacc aaaggtagcg gtaatgacac 1860
agtcctaaat gtggccttgc tgaatgttat ctccaaccag gagtgtaca tcaagcacgc 1920
aggacgtgtg cgggagagtg agatgtgcac tgagggactg ttggcccctg tgggggctg 1980
tgagggtgac tacggggggc cacttgccctg ctttaccac a actgctggg tccctggaagg 2040
aattataatc cccaaccgag tatgcgcaag gtcccgcctg ccagctgtct tcacgcgtgt 2100
ctctgtgttt gtggactgga ttcacaaggt catgagactg ggttagggcc agccttgatg 2160
ccatatgcct tggggaggac aaaacttctt gtcagacata aagccatgtt tctcttta 2219

```

<210> 334

<211> 431

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(431)

<223> 3' terminal sequence. glutathione
s-transferase pi (GSTP1) gene.

<400> 334

```

gaaaggaagg caaactctgc cnccc gctca ggtcccccc aaccctcact gtttcccgtt 60
gccattgatg gggagggttca cgtactcagg ggaggccagg naggcntgna gcttggggccg 120
ggcactgagg cgcacacat atgtgagag cagggggaac gcatccaggc agccagggtc 180
agggaccnca tggatcagca gcaagtccag caggttntag tcagcgaagg agntctggtc 240
tcccacaatg aaggtcttgc ctccctggtt ctgggacagc aggggtctcaa aaggcttcag 300

```

217/292

ttgcccgggc agtgccttca catagtcac cttgcccgcc tcatagttgg tntagatgag 360
 ggagatgtat ttgcagcgga ggtcctccac gccgttcatt tcacctgtcc accagggctg 420
 nctccntttt t 431

<210> 335

<211> 305

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(305)

<223> 5' terminal sequence. glutathione
s-transferase pi (GSTP1) gene.

<400> 335

nattcgccac aggtcgccac catgccgccc tacaccgtgg tctatttccc agttcgagggc 60
 cgctgcctgt cggcaatgct gctggcagat cagggccaga gctggaagga ggaggtngtg 120
 accgtggaga cgtggcagga gggctcactc aaagcctcct gcctatacgg gcagctcccc 180
 aagttccagg acggagacct naccctgtac cagt ccaata ccatcctgcg tcacctgggc 240
 cgcacccttg ggctnctatg ggaaggacca gcaggangca gccctggtgg acatngtgaa 300
 tgacg 305

<210> 336

<211> 737

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(737)

<223> glutathione s-transferase pi (GSTP1) gene.

<400> 336

ggagtttgcg cgcgcagtc ttgccacca tgccgcccta caccgtggtc tatttcccag 60
 ttcgaggccg ctgcgcggcc ctgcgcatgc tgctggcaga tcagg gccag agctggaagg 120
 aggaggtggt gaccgtggag acgtggcagg agggctcact caaagcctcc tgcctatacag 180
 ggcagctccc caagttccag gacggagacc tcaccctgta ccagtccaat accatcctgc 240
 gtcacctggg ccgcaccctt gggctctatg ggaaggacca gcaggaggca gccctggtgg 300
 acatggtgaa tgacggcgtg gag gacctcc gctgcaaata catctccctc atctacacca 360
 actatgaggc gggcaaggat gactatgtga aggcactgcc cgggcaactg aagccttttg 420
 agaccctgct gtcccagaac cagggaggca agaccttcac tgtgggagac cagatctcct 480
 tcgctgacta caacctgctg gacttgctgc tgatccatga ggtcctagcc cctggctgcc 540
 tggatgcgtt cccctgctc tcagcatatg tggggcgctt cagcgcccg cccaagctca 600
 aggccttccct ggctccctt gagtacgtga acctcccat caatggcaac gggaaacagt 660
 gagggttggg gggactctga gggggaggca gagtttgcct tcctttctcc aggaccaata 720
 aaatttctaa gagagct 737

<210> 337

<211> 372

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(372)

<223> 3' terminal sequence. b-cell cll/lymphoma 2
(BCL2) gene.

<400> 337

```
gtgggncgtgt gttgaaacag gccacgtaaa gcaactctct aaaggtcaaa ccaccataga 60
tttgaatctg ctggtcattc gccatctgga tttttaactg aatgaatctc atgggtttta 120
ccaaacatgc atgtaatcct gaataccatg anttaaatgc gganttgccc agggacgagg 180
aaaccttcaa gaaacaaggt caaagggaca ncagatata a ctgtcacant aaacanttct 240
gttgacgtgg gaaatgcaca tgacttggtt gaaacaaagc tcctcagtgg gccagtgaca 300
tccnggggtt ttcttagggt aggctgagga ctcaggggct tatctcacct tctcaggaat 360
gctttttgaa gg                                     372
```

<210> 338

<211> 508

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(508)

<223> 5' terminal sequence. b-cell cll/lymphoma 2
(BCL2) gene.

<400> 338

```
nttcggcaca gacttttttt aagctaccaa ttg tgccgag aaaagcattt tagcaattta 60
tacaatatca tccagtacct taaaccctga ttgtgtatat tcatatattt tggatacgca 120
ccccccaact cccaatactg gctctgtctg agtaagaaac agaatcctct ggaacttgag 180
gaagtgaaca tttcgggtgac ttcogcatca ggaaggctag agttaccagc agcatcaggc 240
cgccacaagt gcctgctttt aggagaccga agtccgcaga acctgcctgt gtcccagctt 300
ggaggcctgg gtctctggaa ctgagccggg gccctcactg gccttccttc caggggatgg 360
atcaacaggg gcagtgtggt cttccgaatg tctgggaagc tgatgggagc tcagantttc 420
cactgtcaag aaagaggcag ttaggagggg tttgggtggg gcttgttcac ctgg ggggcc 480
ttccaggtag ggcccttttt aagtggga                                     508
```

<210> 339

<211> 445

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(445)

<223> 3' terminal sequence. proliferating cell
nuclear antigen (PCNA) gene.

<400> 339

219/292

```

tttttttant caaaagtgtt aaattcaagt aactttattt aaattcaaaa acaattctta 60
aaactgcatt tagagtcagg acccttttgt attataaaaa tcacaagtat ttctaagaga 120
caaaaatact totaggttaa ctagaccaga tctgactt tg gactttattc tttaaacaan 180
ttgcagagan tagagaaaaa antaggttat ttacagaaaa caatatctac atatgtactt 240
ngnggtacaa ntttgggtga cagaaaagac ttcaggtata tgctgggcat cttagggaagn 300
cagttctcaa agggnccttag gttttatttn cttggatttt taaggattgc cctaagganc 360
ccttcttcat cctcgn tctt gggggnggcc aggtaggtn tttagggtgc ccntatccc 420
ganttttata ctctncaccg ggggg                                     445

```

<210> 340

<211> 437

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(437)

<223> 5' terminal sequence. proliferating cell
nuclear antigen (PCNA) gene.

<400> 340

```

gctccagcgt tgtaaacctg cagagatgga ctggtccac gtctcttttg tgcagctcac 60
cctgcggtct gagggcttcg acacctaccg ctgcgaccgc aacctggcca tgggggtgaac 120
ctcaccagta tgtccaaaat actaaaatgc gccggcaatg aagatatcat tacactaagg 180
gccgaagata acgcggatac cttggcgcta gtatttgaag caccaaacca ggagaaagtt 240
tcagactatg aaatgaagtt gatggattta gatgttgaac aacttngaag tccagaacag 300
gagtacagct gtgtagtaaa gatgcontct ggtgaatttg c acgtatatg ccgagatctc 360
agccatattg ggagatgctg ttgtaatttc ctgtgncaaa agacgggagt gaaaattttt 420
ctgcaagtgg gagnact                                     437

```

<210> 341

<211> 1231

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1231)

<223> proliferating cell nuclear antigen (PCNA)
gene.

<400> 341

```

aggctctcagc cggctcgtgc gacgttcgcc cgtctcgtct gaggtctctg aagccgaaac 60
tagctagact ttctctcttc ccgcctgcct gtagcggcgt tgttgccact ccgccaccat 120
gttcgaggcg cgcttggtcc agggctccat cctcaagaag gtgttgagg cactcaagga 180
cctcatcaac gaggcctgct gggatattag ctccagcggg gtaaacctgc agagcatgga 240
ctcgtccac gtctcttttg tgcagctcac cctgcggtct gagggcttcg acacctaccg 300
ctgcgaccgc aacctggc ca tgggcgtgaa cctcaccagt atgtccaaa tactaaaatg 360
gcgcggcaat gaagatatca ttactaaag gccggaagat aacgcggata ccttggcgct 420
agtatgtgaa gcaccaaac aggagaaagt ttcagactat gaaatgaagt tgatggattt 480
agatgttgaa caacttgga ttccagaaca ggagtacagc tgtgtagtaa agatgccttc 540
tggtgaattt gcacgtatat gccgagatct cagccatatt ggagatgctg ttgtaatttc 600
ctgtgcaaaa gacggagtga aattttctgc aagtggagaa cttggaaatg gaaacattaa 660

```

220/292

```

attgtcacag acaagtaatg tcgataaaga ggaggaagct gttaccatag agatgaatga 720
accagttcaa ctaacttttg cactgaggta cctgaact tc ttacaaaag cactccact 780
ctcttcaacg gtgacactca gtatgtctgc agatgtaccc cttgtttag agtataaaat 840
tgcgatatg ggacacttaa aatactactt ggctcccaag atcgaggatg aagaaggatc 900
ttaggcattc ttaaaattca agaaaataaa actaagctct ttgagaactg cttctaagat 960
gccagcatat actgaa gtct tttctgtcac caaatttgta cctctaagta catatgtaga 1020
tattgttttc tgtaaaataac ctattttttt tctctattct ctccaatttg ttaaagaat 1080
aaagtccaaa gtctgatctg gtctagttaa cctagaagta tttttgtctc ttagaaatac 1140
ttgtgatttt tataatacaa aagggtcttg actctaaatg cagttttaag aagtg ttttt 1200
gaatttaaat aaagttactt gaatttcaa c

```

1231

<210> 342
 <211> 383
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(383)
 <223> 3' terminal sequence. adenovirus 5 ela
 binding protein (BS69) gene.

```

<400> 342
tttttttttt aaacacacan gttttcacgc tgtagtaact tggaaatgtg caaccogtgt 60
caacagagac agaaaagcca aagtaacacg aatctcactt tcatgcagct atcagttaaa 120
tattacatac totggaatga ttttacacca aaaatat t c cacaattact tgctctcata 180
ggggtggatc gaagtcttaa aacttgaaaa acaatcaaa aagggttaagt gttctcggtt 240
ctgacatctc catcagcgcc acacactgtg gngaacactg gactaattac acagcaacaa 300
ggaggggaac gatgatgcca agttactgca taatttaggg tacattgtat ggaatggggg 360
gctactgggg gtactttttt tac

```

383

<210> 343
 <211> 483
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(483)
 <223> 5' terminal sequence. adenovirus 5 ela
 binding protein (BS69) gene.

```

<400> 343
gttnaaattg cagggactgg ggtaatcttt tactgagctg gatcttagag aaaatgaata 60
tttaaatttt aaagtttgca catttcatct ttgtcctaac atgagtgtt gtaacaaaat 120
aaacaacaaa aacaaagcca aaaactacct ttatccatat gtgaaattat agatgaggca 180
tacgaatttg tttaatgctt cccttccctt cccacatata atctcactgc ctattatctg 240
gtgtcacctc atgtatcgta agttaatact aaaagaagag aaagcactta agtttcacag 300
aagccgttat gttttaggt aatgggggtc ttgcctaag gaactccatc actgtacaca 360
gaatgaagga nttaatgcca tgtaattttt cttgttattt aagg atgccc tggatttggg 420
aaaaggtctg gtatttttgc gggatgtctg gggtagggga ggccttacct atagggngtg 480
ggg

```

483

221/292

<210> 344
<211> 2722
<212> DNA/RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(2722)
<223> adenovirus 5 ela binding protein (BS69)
gene.

<400> 344
ggagcataat gctaaagaag taaacaggto atggcacgtt taacaaaaag acgacaggcg 60
atacaaaagc tatocagcat ctttgggcag ccattgagat tata cggaac cagaagcaga 120
ttgccaacat tgaccgtatt acaaaatgtg aaacaactac attattcttg aacctatggt 180
gattttttaca tcattacaca gatatgtcat tttcattagt tgtatcattg ttataaactg 240
gtatatgtct cgagtccacg gtatgcaccc taaagagacc acccgtcagc tgagcttagc 300
tgtgaaagat ggtcttattg tc gaaactct aacagtgggc tgcaaagggt caaaagctgg 360
tattgaacaa gaaggatatt ggttgccagg agatgagatt gactgggaaa cagaaaatca 420
tgactgggat tgttttgaat gccatttgcc tggagaggtg ttgatattgtg acctgtgttt 480
tcgtgtgtat cattccaagt gtttgtctga tgagttcagg cttagagaca gcagtagtcc 540
ctggcagtgcc ccagttttgca ggagcattaa gaagaagaat acaaacaaac aggagatggg 600
cacatacctc agattcattg tctcccgcat gaaggagagg gctatagatc ttaataaaaa 660
ggggaaggac aataaacacc cgatgtacag gaggctggtg cactcagctg tggacgttcc 720
caccattcaa gagaaagtga atgaagggaa ataccgaagt ta tgaagagt tcaaagctga 780
tgcccaattg cttctccaca ataccgtgat tttctatgga gcagacagtg agcaagctga 840
cattgcgagg atgctatata aagacacatg tcatgagctg gatgaactgc agctttgcaa 900
gaattgcttt tacttgtcaa atgctcgtcc tgacaactgg ttctgttacc cttgtatacc 960
taatcatgag ctggttttggg ctaaaatgaa aggttttggg ttttggccag ccaaagtcac 1020
gcagaaagaa gacaatcaag tcgacgttcg cttctttggc caccaccacc agagggcctg 1080
gattccttct gaaaacattc aagatatcac agtcaacatt catcggtgc acgtgaagcg 1140
cagtatgggt tggaaaaagg cctgtgatga gctggagctg catcagcgtt tcctacgaga 1200
aggagatatt tggaaatcta agaattgagg ccgaggtgag gaagaggcag aatccagtat 1260
ctcctccacc agtaattgagc agctaaaggt cactcaagaa ccaagagcaa agaaaggacg 1320
acgtaatcaa agtgtggagc caaaaaagga agaaccagag cctgaaacag aagcagtaag 1380
ttctagccag gaaataccca cgatgcctca gccc atcgaa aaagtctccg tgtcaactca 1440
gacaaagaag ttaagtgcct cttcaccaag aatgctgcat cggagcacc agaccacaaa 1500
cgacggcggt tgtcatgaca tgtgccatga caaatacacc aagatcttca atgacttcaa 1560
agaccggatg aagtccgacc acaagcggga gacagagcgt gttgtccgag aagctctgga 1620
gaagctgcgt tctgaaatgg aagaagaaaa gagacaagct gtaaataaag ctgtagccaa 1680
catgcagggt gagatggaca gaaaatgtaa gcaagtaaa gaaaagtgtg aggaggaatt 1740
tgtagaagaa atcaagaagc tggcaacaca gcacaagcaa ctgattttct agaccaagaa 1800
gaagcagtggt tgctacaact gtgaggagga ggccatgtac cactgctg ct ggaacacatc 1860
ctactgctcc atcaagtgcc agcaggagca ctggcacgcg gagcacaagc gcacctgccg 1920
ccggaaga tgaagctggc ccttcccga gtcaccccga tgattactct tttcagacac 1980
agcggttttt gtttccaaga agccaaaatt gtttagaatt tgcttcccat tttgaccag 2040
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ctccgaagtt tttcaggggg taaaagtaac atcagtggag ggtattattt taaataaatt 2160
ttaattgaga atttgttgca ttttcagcaa attttaaaac atttttaggt tttacagaga 2220
ttttaacctt taaacaacag atctttaaaa aacaggtgaa tacaagttag tttacaaaag 2 280
aaacatttag aatagatctg aatgtaagaa ctacagaact gtttcagaaa taaaacatac 2340
taccttgatg tgacattttt ttcttaacct tgttgagctg gttttgttca gcttaattta 2400
ctgttcaaag gcattatctg ttggtcacac cagtgggtat atgattgaat ttagggaaca 2460
gggttgacac agcagggcta gtctgcata tttttt cttt aatatttccc aattgtgttt 2520
ttcattattt cttttcaata tataactttt ataacaaatt attagctttg atctttagt 2580
ttaaaattgc agggaaactg gtaaatcttt tactgagctg gatcttagag aaaatgaata 2640
tttaaatatt aaagtttgc acattttcac tttgtcctaa catgagtgtc tgaacaaaa 2700

taaaacaaca aaaacaaagc ct

2722

<210> 345

<211> 363

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(363)

<223> 3' terminal sequence. matrix
metalloproteinase 11 (stromelysin 3) (MMP11) gene.

<400> 345

```
gcatgcagca tcctgagtgg tagcgtcgat ctacagagggc acccctctcc agtcagtggc 60
cctgcgggna cggngactgt ctacacgccg ggtgctgggg tggaaacgcc agtagtccct 120
gcctcggaag aagtagatct tggtcttctc gggacccag accaaggc ag catggaccgg 180
gaacctcacc agggcccagc tcggtgaggg gtgcggggcc cagggactgg cttttcaccg 240
tcgtacacc agtacttgag caccttgga agaaccaaat gtgggcccg cttaccagc 300
attggccttt tcgccacagg gctggggcag tccctgccag tngcgagaag ccaattttgg 360
gca 363
```

<210> 346

<211> 2260

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2260)

<223> matrix metalloproteinase 11 (stromelysin 3)
(MMP11) gene.

<400> 346

```
aagcccagca gcccggggc ggatggctcc ggccgcctgg ctccgcagcg cggccgcgcg 60
cgccctctctg ccccgatgc tgctgctgct gctccagccg ccgccgctgc tggccggggc 120
tctgccgcgg gagtccacc acctccatgc cgagaggagg gggccacagc cctggcatgc 180
agccctgccc agtagcccg cacctgcccc tgccacgcag gaagccccc ggcctgccag 240
cagcctcagg cctccccgt gtggcgtgcc cgacctct gatgggctga gtgccgcaa 300
ccgacagaag aggttcgtgc tttctggcg gcgctgggag aagacggacc tcacctacag 360
gatccttcgg ttcccatggc agttggtgca ggagcaggtg cggcagacga tggcaga ggc 420
cctaaaggta tggagcgatg tgacgccact cacctttact gaggtgcacg agggccgtgc 480
tgacatcatg atcgacttcg ccaggtactg gcattggggac gacctgccgt ttgatgggccc 540
tgggggcac cttggccatg cttcttccc caagactcac cgagaagggg atgtccactt 600
cgactatgat gagacctgga ctatcgggga tgacc agggc acagacctgc tgcaggtggc 660
agcccatgaa tttggccacg tgctggggct gcagcacaca acagcagcca aggcctgat 720
gtccgccttc tacacctttc gctacccact gactctcagc ccagatgact gcaggggcgt 780
tcaaacctta tatggccagc cctggccac tgtcacctcc aggacccag ccttggggccc 840
ccaggctggg atagacacca atgagattgc accgctggag ccagacgccc cgccagatgc 900
ctgtgaggcc tcctttgacg cgtctccac catccgaggc gagctctttt tcttcaaagc 960
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tcgccactgg cagggactgc ccagccctgt ggacgctgcc ttcgaggatg ccca gggcca 1080
catttggttc ttccaagggtg ctcagtactg ggtgtacgac ggtgaaaagc cagtcctggg 1140
```

223/292

```

ccccgcaccc ctcaccgagc tgggcctggt gaggttcccg gtccatgctg ccttgggtctg 1200
gggtcccagag aagaacaaga tctacttctt ccgaggcagg gactactggc gtttccacccc 1260
cagcaccocgg cgtgtagaca gtcccgtagc cccgaggggcc actgactgga gaggggtgcc 1320
ctctgagatc gacgctgcct tccaggatgc tgatggctat gcctacttcc tgcgcggccg 1380
cctctactgg aagtttgacc ctgtgaaggt gaaggctctg gaaggcttcc cccgtctcgt 1440
gggtccctgac ttctttggct gtgccgagcc tgccaacact ttctctgac catggcttgg 1500
atgccctcag ggggtgctgac ccctgccagg ccacgaatat caggctagag acccatggcc 1560
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tgagcaactg ggctgtaagg caggggccact tcctgaggtc aggtcttggg aggtgcctgc 1980
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caacatacct caatcctgtc ccaggccgga tcctcctgaa gcccttttgc cagcac tgct 2160
atcctccaaa gccattgtaa atgtgtgtac agtgtgtata aaccttcttc ttcttttttt 2220
tttttaaaact gaggattgtc attaaacaca gttgttttct 2260

```

<210> 347

<211> 273

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(273)

<223> 3' terminal sequence. hypothetical protein
mgl13071 (MGC13071) gene.

<400> 347

```

atgtttattg aacgtaacag tatatttcat gtagtttccc ataatttttt catgtactaa 60
ctcatgtaat tctttgtttt ttagagatct gaagtgat tt tacctttact tccttcaactt 120
taagccaatc atgaaatttc agtgatttct ggggtgaggg cgaaagggtg tgttacgaat 180
catcggggct gtggccagnt tgcctcacgg aggtgcaggt aggtcggggc ctactaggg 240
canctggagg agcacggact gccctgccgg cag 273

```

<210> 348

<211> 330

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(330)

<223> 5' terminal sequence. hypothetical protein
mgl13071 (MGC13071) gene.

<400> 348

```

ggagtagaca acattgtggt aggggaaggg actcactttc tcattcccatg tgtacaaaaa 60
ccaattatct ttgactgctg ttctcaacca cgtagtcgnc cagtcacac tggtagcaaa 120
gatttagaca atgtcaacat cacactgtgc atcctcttcc ggcccatcac tagccagctt 180

```

224/292

```

cctcgcatct tcaccagcat tggagaggac tacgatgagt gtgtgctgcc gttcattacc 240
acggagatcc tcaagtcact ggtggctcgc tttgatgctg gagaactaat caccagagg 300
gagcttggtt tccagcngg tgaagnacca                               330

```

<210> 349

<211> 1168

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:prime r

<220>

<221> misc_feature

<222> (1)..(1168)

<223> hypothetical protein mgc13071 (MGC13071)
gene.

<400> 349

```

aaatgatgat agtagtacct acagtatagt gctgttagaa ttacatgagt tagatgtgga 60
ggtcagagtg gaagcagggtg tgagaggggtc ccgcagaaga aaacatggct gccaaagtgt 120
ttgagtccat cggcaagttt ggccctggcct tagctgttgc aggaggcatg gtgacctctg 180
ccttatgtaa tgtggatgct gggcacagag ctgccatctt tgaccaattc cgtggagtac 240
agaacattgt ggtaggggaa gggactcact ttctcatccc atgtgtacaa aaaccaatta 300
tctttgactg ctgttctcaa ccacgtagtg cgccagtcac c actggtagc aaagatttac 360
agaatgtcaa catcacactg tgcacacctt tccggcccat cactagccag ctctctcgca 420
tcttcaccag cattggagag gactacgatg agtgtgtgct gccgttcatt accacggaga 480
tcctcaagtc actgggtggct cgctttgatg ctggagaact aatcaccagc agggagctgg 540
tctccagcca ggtgagcaac aaccttatgg agtgagcagc cacctttggg ctcatctctg 600
acgacgtgtc tttagacacat ctgaccttct tgaaggagtt aacagacagg tggcccccca 660
ggaagcagag agcgccagat ttgtggtgga aaaggcggcc atcatctctg ctgagggtga 720
ctccaaggca gctgagctga tcgccaactc actggccact gcaggggacg gccagagcga 780
gctgtgcaag ctggaagctg cagaagacat tgcataccag ctctcatgct ctcggaacat 840
cacctgcctg ccggcagggc agtccgtgct cctccagctg ccctagttag gccccagcct 900
acctgcacct ccgtgaggca actggggccac agccccgatg attcgtaaca ccacctttcg 960
ccctcacccc agaaatcact gaaatttcat gattggctta aagtgaagga agtaaaggta 1020
aaatcacctc agatctctaa aaaacaaaga attacatgag ttagtacatg aaaaaattat 1080
gggaaactac atgaaatata ctgttacgtt caataaacat tagcttctgt atataaaaaa 1140
aaaaaaaaa aaaaaaaaaa aaaaaaaa                               1168

```

<210> 350

<211> 315

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(315)

<223> 5' terminal sequence. interleukin enhancer
binding factor 2, 45kd (ILF2) gene.

<400> 350

```

ctggctttga aatcagttct ncgtgatgct acagtgaaga ttctcattac aacagtgcc 60
cccaatcttc gaaaactgga tccagaactc catttgata tcaaagtatt gcagagtgc 120
ttagcagcca tccgacatgc ccgtgggttc gaggaaaatg cttctcagtc cacagttaaa 180
gttcnccanc agantactga aggacttgag gattcgtttt ccnggctttg agc cctcaca 240

```

225/292

cccnggatnc ttgaactact aggn cattat gctgtgatga acaacccac caganagcct 300
 ttggcnctaa acgtt 315

<210> 351

<211> 1552

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1552)

<223> interleukin enhancer binding factor 2, 45kd
 (ILF2) gene.

<400> 351

cggttggtgc ggcctccatt gtctgtgttt taaggcgcca tgaggggtga cagaggccgt 60
 ggtcgtggtg ggcgcttttg ttccagagga ggcccaggag gag ggttcag gccctttgta 120
 ccacatatcc catttgactt ctatttgtgt gaaatggcct ttccccgggt caagccagca 180
 cctgatgaaa ctctcttcag tgaggccttg ctgaagagga atcaggacct ggctcccaat 240
 tctgtgaac aggcacttat cctttctctg gtgacaaaaa taaacaatgt gattgataat 300
 ctgattgttg ctccagggaac a ttggaagt gaaattgaag aagttcgaca ggtgggatcc 360
 tataaaaagg ggacaatgac tacaggacac aatgtggctg acctgggtgt gatactcaag 420
 attctgcca cgttggaagc tgttgctgcc ctggggaaca aagtcgtgga aagcctaaga 480
 gcacaggatc ctctgaagt tttaaccatg ctgaccaacg aaactggctt tgaaatcagt 540
 tcttctgatg ctacagtga gattctcatt acaacagtgc cacccaatct tcgaaaactg 600
 gatccagaac tccatttggg tatcaaagta ttgcagagt gcttagcagc catccgacat 660
 gcccgctggt tcgaggaaaa tgccttctcag tccacagtta aagttctcat cagactactg 720
 aaggacttga ggattcgctt tccctggctt gagccctca c acctggat ccttgacct 780
 ctaggccatt atgctgtgat gaacaacccc accagacagc ctttgccct aaacgttgca 840
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 actgaccct gtgagagtgg caactttaga gtacacacag tcatgacct agaacagcag 960
 gacatggtct gctatacagc tcagactctc gtccgaatcc tctcacatgg tggctttagg 1020
 aagatccttg gccaggagg tgatgccagc tatcttgct ctgaaatata tacctgggat 1080
 ggagtgatag taacaccttc agaaaaggct tatgagaagc caccagagaa gaaggaagga 1140
 gaggaagaag aggagaatac agaaagaacc acctcaagga gaggaagaag aaagcatgg a 1200
 aactcaggag tgacattccc ttcactcctt ttctaccca agggaaagac tggagcctaa 1260
 gctgcctgt actggcttta catggtgaca gacattcgt gcataggaag atagcaggag 1320
 aaagtaactc catagagtgt cattccactg gttgatattg gcttagctgc cagtctccca 1380
 tttgtacct atgccatcca tctataatgg agg ataccaa cattctctcc taatattcta 1440
 taatctccaa ctctgaaaaa cccctctctc aactaatact ttgctgttga aatgttgtga 1500
 aatgttaagt gtctggaaat tttttttct aagaaaaact attaaagtac tt 1552

<210> 352

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(396)

<223> 3' terminal sequence. hypothetical protein
 flj11307 (FLJ11307) gene.

<400> 352

ctccattaca ggggtttttgc cacttgetgt gaggataggg ccctgagttc ttacctctaa 60
 ggtactggag gtttcagttg tagaatttcc agtattattg cttgagtttg aagacactgt 120
 ttcattttta ctttcattat ctgatttttc atcggaaactc atacattcaa tatctgcac 180
 aaagcctgtt ggatatccca ttgectgcaa taccttcacc gctacgtgaa agttttgctg 240
 tttcttggga tggctctgag gcttcataatg ttgtgccaac cacatctaca gacattg tga 300
 agactggggg catgaacggg ggccagactg aagataagaa gctatactga agcacaggcc 360
 tgatctgaat taagctcat tagtgccatt ccataa 396

<210> 353

<211> 1858

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer

<220>

<221> misc_feature

<222> (1)..(1858)

<223> hypothetical protein flj11307 (FLJ11307)
gene.

<400> 353

tcgatgaaag atcctccgga cttattggac aggcagaaat gcccgaaagc cttggcgtct 60
 cttcgacatg ccaaatgggt tcaggcaagg gcaaatggat taaatcatg tgtaat tgtc 120
 ctccgcattc tgcgtgattt gtgcaacaga gtccccacat gggcaccatt gaaaggatgg 180
 ccactagaac ttatatgtga aaagtctata ggtacttgta atagacctt gggcgtggg 240
 gaggccttga gacgagtaat ggagtgttg gcatctggaa tactaottec tgggggtcct 300
 ggtcttcattg atccttgtga gcgagacca acag atgctc tgagctatat gaccatccag 360
 caaaaagaag atattaccca cagtgcacag catgcactca gactatcagc ctttgccag 420
 atttacaag tgctggagat ggacccctt ccactagta agccttttca gaagtattcc 480
 tggctcagtt ctgataaaga aggtgctggg tcttcagctc taaagaggcc atttgaagat 540
 ggattagggg atgataaaga cccaacaag aagatgaaac gaaacttaag gaaaattctg 600
 gatagtaaag caatagacct tatgaatgca ctaatgaggc taaatcagat caggcctggg 660
 cttcagtata agctcctatc tcagtctggc cccgttcatt cccagctctt cacaatgtct 720
 gtgatgttg atggcacaac atatgaagcc tcaggaccat ccaagaaaac agca aaactt 780
 cacgtagcgg tgaaggattt gcaggcaatg ggatatccaa caggctttga tgcagatatt 840
 gaatgtatga gttccgatga aaaatcagat aatgaaagta aaaatgaaac agtgtcttca 900
 aactcaagca ataatactgg aaattctaca actgaaacct ccagtacctt agaggtaaga 960
 actcagggcc ctatcctcac agcaagtggc aa aaaccctg taatggagct caatgaaaaa 1020
 agaagaggtc tcaagtatga actcatctca gagactgggt gaagccatga caagcgcttt 1080
 gtaatggagg tagaagtaga tggacagaaa ttcagaggcg caggctccaa taagaaagt 1140
 gcaaaggcga gtgcagcttt agctgccttg gagaaactgt tttctggacc caatgcggca 1200
 aataataaga aaaagaagat tatcctcag gcaaagggcg ttgtgaatac agctgtgtct 1260
 gcagcagtc aagctgttcg gggcagagga agaggaaact taacaagggg agctttttgt 1320
 ggggcgacag ctgctcctgg ctacatagct ccaggctatg gaacaccata tggttacagc 1380
 acagctgccc ctgcctatgg tttaccaag agaattgttc tgttac ccgt tatgaaattt 1440
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 tgaacaacaa tacagtacaa cacagaatgt tagagaaaa gcctttttat cctgctttct 1560
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 acggtacctc atttctggaa aataacttat accagccctt ctgttctagg gaaataaaag 1740
 tctagcagtt caaagtttaa gttttaagag acgtatcaga ttatgtaaaa ttaaatttgt 1800
 gaaggatgta tagagtctca aacactgac acaataaac tgctttgttg taacacag 1858

<210> 354

<211> 242

<212> DNA

227/292

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(242)

<223> 5' terminal sequence. v-myb avian
myeloblastosis viral oncogene homolog (MYB) gen e.

<400> 354

```

agaaccccag ctatcaaaag gtcaatctta gaaagctctc caagaactcc tacaccattc 60
aaacatgcac ttgcagctca agaaattaaa tacggtcccc tgaagatgct acctcagaca 120
ccctctcatc tagtagaaga tctgcaggat gtgatcaaac aggaatctga tgaatctgga 180
attgttgctg agtttcaacg aaaatggacc acccttactg aacgaaaatc ntacaacgag 240
gt                                     242

```

<210> 355

<211> 3225

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(3225)

<223> v-myb avian myeloblastosis viral oncogene
homolog (MYB) gene.

<400> 355

```

ggcggcagcg ccctgccgac gccggggagg gacgcaggca ggcggcgggc agcggggaggc 60
ggcaccgccg tgctccccgc ggctctcggc ggagccccgc cgcccgccgc gccatggccc 120
gaagaccccg gcacagcata tatagcagtg acgaggatga tgaggacttt gagatgtgtg 180
accatgacta tgatgggctg cttcccaagt ctggaaagcg tcaattgggg aaaacaagggt 240
ggacccggga agaggatgaa aaactgaaga agctggtgga acagaatgga acagatgact 300
ggaaagttaa tgccaattat ctcccgaatc gaacagatgt gcagtgccag caccgat ggc 360
agaaagtact aaaccctgag ctcatcaagg gtccttggac caaagaagaa gatcagagag 420
tgatagagct tgtacagaaa tacggtccga aacgttggtc tgttattgcc aagcacttaa 480
aggggagaat tggaaaacaa tgtagggaga ggtggcataa ccaattgaat ccagaagtta 540
agaaaacctc ctggacagaa gaggaagaca gaatt atta ccaggcacac aagagactgg 600
ggaacagatg ggcagaaatc gcaaagctac tgccctggacg aactgataat gctatcaaga 660
accactggaa ttctacaatg cgtcggaagg tcgaacagga aggttatctg caggagtctt 720
caaaagccag ccagccagca gtggccacaa gcttcagaa gaacagtcac ttgatgggtt 780
ttgctcagc tccgctaca gctcaactcc ctgcccactg ccagcccact gtaacaacg 840
actattccta ttaccacatt tctgaagcac aaaatgtctc cagtcatggt ccataacctg 900
tagcgttaca tgtaaatata gtcaatgtcc ctccagccagc tgccgcagcc attcagagac 960
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tgtcaaccga gaatgagcta aaaggacagc aggtgctacc aacacagAAC cacacatgca 1080
gtacccccgg gtggcacagc accaccattg ccgaccacac cagacctcat ggagacagtg 1140
cacctgtttc ctgtttggga gaacaccact ccaactccatc tctgccagcg gatcctggct 1200
ccctacctga agaaagcgcc tcgccagcaa ggtgcatgat cgtccaccag ggcaccattc 1260
tggataatgt taagaacctc ttagaatttg cagaaacact ccaatttata gattctttct 1320
taaacacttc cagtaaccat gaaaactcag acttggaat gccttcttta acttcaccc 1380
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ctcaaaagga taataactgtt tttagaacct cagctatcaa aaggtcaatc ttagaaagct 1500
ctccaagaac tcctacacca ttcaaaccat cacttgacgc tcaagaaatt aaatacggtc 1560
ccctgaagat gctacctcag acacctctc atctagtaga agatctgcag gatgtgatca 1620

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228/292

```

aacaggaatc tgatgaatct ggatttgttg ctgagtttca aga aaatgga ccacccttac 1680
tgaagaaaat caaacaagag gtggaatctc caactgataa atcaggaaac ttcttctgct 1740
cacaccactg ggaaggggac agtctgaata cccaactgtt cacgcagacc tcgcctgtgc 1800
gagatgcacc gaataattctt acaagctccg ttttaatggc accagcatca gaagatgaag 1860
acaatgttct caaagcat tt acagtaccta aaaacaggtc cctggcgagc cccttgacgc 1920
cttgtagcag tacctgggaa cctgcatcct gtggaagat ggaggagcag atgacatctt 1980
ccagtcaagc tcgtaaatac gtgaatgcat tctcagcccg gacgttggtc atgtgagaca 2040
tttcagaaa agcattatgg ttttcagaac agttcaagtt gacttgggat atatcat tcc 2100
tcaacatgaa acttttcatg aatgggagaa gaacctatct ttgttgtggt acaacagttg 2160
agagcacgac caagtgcatt tagttgaatg aagtcttctt ggatttcacc caactaaaag 2220
gatttttaaa aataaataac agtcttacct aaattattag gtaatgaatt gtagccagtt 2280
gttaatatct taatgcagat ttttttaaaa aaaaacataa aatgatttat ctggtatttt 2340
aaaggatcca acagatcagt attttttctt gtgatgggtt ttttgaaatt tgacacatta 2400
aaaggatcca cagtatttca cttttctcga tcactaaaca tatgcataata tttttaaaaa 2460
tcagtaaaaag cattactcta agtgtagact taataccatg tgacatttaa tccagattgt 2520
aaatgctcat ttatggttaa tgacattgaa ggtacattta ttgtacaaaa ccattttatg 2580
agttttctgt tagcttgcct taaaaattat tactgtaaga aatagtttta taaaaaatta 2640
tatttttatt cagtaattta attttgtaaa tgccaaatga aaaacgtttt ttgctgctat 2700
ggcttagacc ttagacatg ctgctagtat cagaggggca gtaga gcttg gacagaaaga 2760
aaagaaactt ggtgttaggt aattgactat gcactagtat ttcagacttt ttaattttat 2820
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attttttatt tgggtttttt tgttattgtt ggtttatata agcatgcgtt gcacttcttt 2940
tttgggagat gtgtgttgtt catgttctat gttttgtttt gtgtgtagcc tgactgtttt 3000
ataatttggg agttctcgat ttgatccgca tccctgtgg tttctaagt tatgttctca 3060
gaactgttgc atggatcctg tgtttgcaac tggggagaca gaaactgtgg ttgatagcca 3120
gtcactgcct taagaacatt tgatgcaaga tggccagcac tgaacttttg agatatgac g 3180
gtgtacttac tgccttgtag caaaataaag atgtgccctt atttt 3225

```

<210> 356

<211> 369

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(369)

<223> 3' terminal sequence. zinc finger protein 9
(a cellular retroviral nucleic acid binding
protein) (ZNF9) gene.

<400> 356

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gtagttaaat gcagaaagtc ggtttttttc cacccttttc ctctttttac acggcaagta 60
aagctcactg gctggggagt tgcctctatc tgccaacctt tggccagtga agaggattca 1 20
gagaaaataa tacaaccatc aatcagaaaa aggaggggag acaaaggaaa ataattagtc 180
ttagctcaa ttgtgcattc ccgtgcaagg tgccctgact cgccacagcg gtaacagttg 240
acttcacttg tcttctgca gttgatggtt acatgaccag tttcaccaca cctatagcac 300
ttcacttttg tgcagtcttt tttgaatgtg tcccgaattc tcccacaaga atancctttc 360
tgctcanct 369

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<210> 357

<211> 1500

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(1500)
 <223> zinc finger protein 9 (a cellular retroviral
 nucleic acid binding protein) (ZNF9) gene.

<400> 357
 gaattccaaa cagcctctac cttgcgagcc gtcttcccca ggctgcgctc cgagtctccg 60
 ccgctgcggg cccgctccga cgcggaagat ctgactgcag ccatgagcag caatgagtgc 120
 ttcaagtgtg gacgatctgg ccactgggcc cgggaatgtc ctactgggtg aggcctgtgt 180
 cgtggaatga gaagccgtgg cagaggtggt ttacctcgg atagaggttt ccagtttgtt 240
 tcctcgtctc ttccagatat ttgttatcgc tgtggtgagt ctggatcatc tgccaaggat 300
 tgtgatcttc aggaggatgc ctgctataac tgcggtagag gtg gccacat tgccaaggac 360
 tgcaaggagc ccaagagaga gcgagagcaa tgcgtctaca actgtggcaa accaggccat 420
 ctggctcgtg actgcgacca tgcagatgag cagaaatgct attcttgttg agaattcggg 480
 cacattcaaa aagactgcac caaagtgaag tgctataggt gtggtgaaac tggatcatgt 540
 gccatcaact gcagcaagac aagtgaagtc aactgttacc gctgtggcga gtcagggcac 600
 cttgcacggg aatgcacaat tgaggctaca gcctaattat ttccctttgt cgcctctcct 660
 tttctgtatt gatggttgta ttatctctc tgaatcctc tcaactggcca aaggttggtg 720
 gatagaggca actcccaggc cagtgcgctt tacttgccgt gtaaaaggag gaaaggggtg 780
 gaaaaaacc gactttctgc atttaactac aaaaaagtt tatgtttagt ttggttagag 840
 tgttatgtat aatgctttgt taaagaacc cctttcgtg ccactggtga atagggtttg 900
 atgaatggga agagttgagt cagaccagta agcccgctc gggttccttg aacatgttcc 960
 catgtaggag gtaaaaccaa ttctggaagt gtctatgaac tccataaat aactttaatt 1020
 ttagtataat gatggtcttg gattgtctga cctcagtagc tattaataaa catcaagtaa 1080
 catctgtatc aggccttaca tagaacatac agttgagtgg gagtaaacaa aaagataaac 1140
 atgcgtgtta atggctgttc gagagaaatc ggaataaaag cctaaacagg aacaacttca 1200
 tcacagtgtt gatgttggac acatagatgg tgatggcaaa ggtttagaac acattatatt 1260
 caaagactaa atctaaaacc cagagtaaac atcaatgctc agagttagca taatttgtag 1320
 ctattcagga attgcagaga aatgcatttt cacagaaatc aagatgttat tttgtatatac 1380
 tatatcactt agacaactgt gtttcatttg ctgtaatcag tttttaaaag tcaga tggaa 1440
 agagcaactg aagtcctaga aaatagaaat gtaattttta actattccaa taaagctgga 1500

<210> 358
 <211> 425
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(425)
 <223> 3' terminal sequence. camp responsive
 element modulator (CREM) gene.

<400> 358
 ttttttactt ctgcaagatc ttttatatta cacagtagag ttaaaaactg tagtaaattg 60
 tcagatattt aaatgagcac caaacactac aaagtgcac caacatgggt ctattaaaaa 120
 ctncctttga ctatggcatt caaggacagc aatacaat ct tttttttttt taacaaagca 180
 actaatataa aaatctgcaa atgccatata ttcatatcta ggotattctt cncatatagg 240
 catgtcatta gatagacttt cttctatctc ttccngagg natTTTTTTT nggtttacnt 300
 ttattgnact gctggatgca ttatttttga tcatcctttc ctaaaatgnt ttaaagacct 360
 gcaataaatt ttattgcata ggacacnatt ggtgncacat agaatgggag cngcaagtat 420
 gtggc 425

<210> 359

<211> 232
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(232)
 <223> 5' terminal sequence. camp responsive
 element modulator (CREM) gene.

<400> 359
 ggatttagag ttaactagct caccactgcc totgcctcca agctgccttt tagactgaat 60
 agcttttctt gttagcccta ctttaacatt tcttttgaag tgggtgtctg cttgaagagg 120
 gaaacacgtc atgaaactgt aatgcatgaa cagaactcag gagttgtctg gccagcttag 180
 tgctgccact ggtgacatgc caacttacca gatccgagct cctantgnng ct 232

<210> 360
 <211> 1431
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:prime r

<220>
 <221> misc_feature
 <222> (1)..(1431)
 <223> camp responsive element modulator (CREM)
 gene.

<400> 360
 atgaccatgg aaacagttga atcccagcat gatggaagta taacagcttc tttgacagag 60
 agcaagtctg ctcatgtgca gactcagact gggcaaattt caatccctgc tttagctcag 120
 tgcagtggag tgagatcagg caccagaaga ggtcccccag ctgtaactct agtgcagtta 180
 ccttcggggc aaactataca tgtccaggga gtaattcaga caccacagcc atgggttatt 240
 cagtcacatcag aaatacacac cgttcaggta gcagcaattg cagagacaga tgaatctgca 300
 gaatcagaag gtgtaattga ttctcataaa cgtagagaaa t cctttcacg aagaccctct 360
 tataggaaaa tactgaatga actgtcctct gatgtgcctg gtgttcccaa gattgaagaa 420
 gagagatcag aggaagaagg aacaccacct agtattgcta ccatggcagt accaactagc 480
 atatatcaga ctagcacggg gcaatacatt gctatagccc aaggtggaac aatccagatt 540
 tctaaccag gatctgatgg tgttcaggga ctgcaggcat taacaatgac aaattcagga 600
 gtcctccac caggtgctac aattgtacag tacgcagcac aatcagctga tggcacacag 660
 cagttctttg tcccaggcag ccaggttgtt gttcaagctg ccactggtga catgccaaact 720
 taccagatcc gagctcctac tgcctgtttg ccacaggagag tggatgatgg tgcacgccc 780
 ggaagtgtgc acagtcccca gcagctggca gaagaagcaa cacgcaaagc agagctgagg 840
 ctaatgaaaa acagagaagc tgcccgggag tgtcgcagga agaagaaaga atatgtcaa 900
 tgtcttgaaa atcgtgtggc tgtgcttgaa aaccaaaca agactctcat tgaggaactc 960
 aaggccctca aagatcttta ttgccataaa gtagagtaac tgtctttgac ttggaccttg 1020
 tttactctaa tcaaggcagg agatgcagca gtccacttta ttgccatgtg gacttgtggg 1080
 aaggacacgt gtgacctta agaatccagt ttggattagt gtttgaaatt gaattgggaa 1140
 tgtgttcca ggaatgggaa tgcagcgtga tcacacttac cgagcttact ttgatctgtt 1200
 tgtcaatagc atgcaaaaaa tgcctttgtt gccctttgct tctgcttttt ttcagggaag 1260
 ctgccaaaga atgtcgacgt cgaaagaaag aatatgtaaa atgcctggag agccgagttg 1320
 cagtgtgga agtcacagac aagaagctta tagaggaact tgaaccttg aaagacattt 1380
 gttctcccaa aactgattac tagaaatatt taactatgaa ctgattacag a 1431

<210> 361
<211> 457
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(457)
<223> 3' terminal sequence. cathepsin b (CTSB)
gene.

<400> 361
caagttggag aaacctttta ttggcacagg cattccttgt taacttgaca gggatgaagct 60
gtaatttttc aaaaacagta aaagctgggt tctcctaaac tattttcctt gtggtagtag 120
agatcagtgg gtcagaaaca actcctgacc acttggttcc cttttgagcc gcgtcattag 180
gaggcaatct gtaaaactag cacaggtctc ccgctgttcc actgggtcac ccacatg att 240
agcagagtgc acgaaaaaat aaaacttcta ttaaagaatc atgctgagca caacatcaga 300
gaggttgatg cattgcaaac tcgatagatg cagggggcct gggagactgg cgttctccaa 360
agggctccca acaccatctc tctctgatt tctgtgacaa atgtggaagc tacttgcttg 420
gaggtactgg gggaactgat ggggaactt tcac cg 457

<210> 362
<211> 401
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(401)
<223> 3' terminal sequence. melan -a (MLANA) gene.

<400> 362
atcatgcatt gcaacattta ttgatggagt ttccccaatt taatatttct catcatttcc 60
tcacatgatt agtactgcta gcggacctac taaaatttta aactgactt attattagag 120
atggttgca tttttcctac accattccaa aggagaacat tagatgtctg tattaattc 180
aagcaaaaagt gtgagagaaa taatttcagc atgtctcagg tgtctcgct g gcncttaagg 240
tgaataagggt ggtggtgact gttctgcaga gagtttctca taagcagggt gagcattggg 300
aaccacaggt tcacagtttt tctcttgaag agacactttg ctgtcccgat gatcaaacc 360
ttcttggtgg catcttcctg ttaaggcaca ttgaggccaa c 401

<210> 363
<211> 370
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(370)
<223> 5' terminal sequence. melan -a (MLANA) gene.

232/292

<400> 363
 attaaggaag gtgtcctgtg ccctgaccct acaagatgcc aagagaagat gctcacttca 6 0
 tctatggtta ccccaagaag gggcacggcc actcttacac caccgctgaa gaggccgctg 120
 ggatcgcat cctgacagt atcctgggag tcttactgct catcggtgt tggatttga 180
 gaagacgaaa tggatacaga gccttgatgg ataaaagtct tcatgttggc actcaatgtg 240
 cttacaaga agatgcccac aagaagggtt tgatcatcgg gacagcaaag tgtctcttca 300
 agagaaaaac tgtgaacctg tggttcccaa tgctccacct gcttatggag aaactctctg 360
 cagaacagtc 370

<210> 364

<211> 1524

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1524)

<223> melan-a (MLANA) gene.

<400> 364
 agcagacaga ggactctcat taaggaaggt gtctgtgcc ctgaccctac aagatgccaa 60
 gagaagatgc tcaacttcac tatggttacc ccaagaagg gacggccac tcttacacca 120
 cggctgaaga ggcgctggg atcggcatcc tgacagtgat cctgggagtc ttactgctca 180
 tcggctggtg gtattgtaga agacgaaatg gatacagagc cttgatggat aaaagtcttc 240
 atgttggcac tcaatgtgcc ttaacaagaa gatgcccaca agaagggtt gatcatcggg 300
 acagcaaagt gtctcttcaa gagaaaaact gtgaacctgt ggttccc aat gctccacctg 360
 cttatgagaa actctctgca gaacagtcac caccacctta tcaccttaa gagccagcga 420
 gacacctgag acatgctgaa attatttctc tcacactttt gcttgaattt aatacagaca 480
 tctaattgtt tcctttggaa tgggttagga aaaatgcaag ccatctctaa taataagtca 540
 gtgttaaaat tttagtaggt ccgct agcag tactaatcat gtgaggaaat gatgagaaat 600
 attaaattgg gaaaactoca tcaataaatg ttgcaatgca tgatactatc tgtgccagag 660
 gtaatgttag taaatccatg gtgttatttt ctgagagaca gaattcaagt gggatttctg 720
 gggccatcca atttctcttt acttgaaatt tggctaataa caaacatagtc aggttttctga 780
 accttgaccg acatgaactg tacacagaat tgttccagta ctatggagtg ctacaaaagg 840
 atacttttac aggttaagac aaagggttga ctggcctatt tatctgatca agaacatgtc 900
 agcaatgtct ctttgtgctc taaaattcta ttatactaca ataatatatt gtaaagatcc 960
 tatagctctt tttttttgag atggagtttc gcttttgttg cccag gctgg agtgcaatgg 1020
 cgcgatcttg gctcaccata acctccgct cccaggttca agcaattctc ctgccttagc 1080
 ctctgagta gctgggatta caggcgtgcg ccactatgcc tgactaattt ttagttttta 1140
 gtagagacgg ggtttctcca tgttggtcag gctggtctca aactcctgac ctacaggtgat 1200
 ctgcccgcct cagcctccca aagtgtctga attacaggcg tgagccacca cgctggctg 1260
 gatcctatat cttaggtaag acatataacg cagtctaatt acatttcaact tcaaggctca 1320
 atgtattctt actaatgac aagtatttct tactaaacca gaaatttgta gaaggattta 1380
 aataagtaaa agctactatg tactgcctta gtgctgatgc ctgtgtactg ccttaaatg t 1440
 acctatggca atttagctct cttgggttcc caaatccctc tcacaagaat gtgcagaaga 1500
 aatcataaag gatcagagat tctg 1524

<210> 365

<211> 556

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

233/292

<221> misc_feature
<222> (1)..(556)
<223> 3' terminal sequence. apr -1 protein (APR-1)
gene.

<400> 365
actattcggtt aggccttttat ttttctctat gttctgcagt aactaaggaa aatcatggta 60
aatgtcaatc ttcacacaac agcagacaca aagggtttca gaaacgtcag atatgaag aa 120
atcctccatc cttcttcaac attttactgg gtatttcaac ttcaaaagaa cagcttattt 180
ctataagtgc tgtacaagat catagattat gatggaacga cttcatttta gaacgttagc 240
aaaactgtta tactaaatgt caatgacagg aaacaaagaa aaaaatttgt tcaattatat 300
ttttaaacat attgttattc tcaacaaacg gaattt taaa acgaatacaa ttttccatta 360
tcaaaaagca aacactctat ttgcgcagttg aacaatgatc actgatacaca aatatacnaat 420
acagtgtccc ccgcccccaa tgcacatcat tttccactta gggaccctgg catccactcc 480
ctgggggtac ccgtgactcc ncctttacac cccccagggg ctggcctcag atctacctaa 540
ggggngggat aacc cc 556

<210> 366
<211> 464
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(464)
<223> 5' terminal sequence. apr -1 protein (APR-1)
gene.

<400> 366
aacagcgcca aggaagctct ggtctggaaa gtgctgggga agttaggaat gcagcctgga 60
cgtcagcaca gcattctttg agatccgaag aagatcgta cagaagagtt tgtgcgcaga 120
gggtacctga tttataaacc ggtgccccgt agcagtcagg tggagtatga gttcttcttg 180
gggccccgag cacacgtgga atcgagcaaa ctgaaagtca tgcattttgt ggcaagggtt 240
cgtaaccgat gctctaaaga ctggccttgt aattatgact gggattcgga cgatgatgca 300
gagggttagg ctatcctcaa ttcaggtgct aggggttatt ccgcccctta agtagatctg 360
gaggcagacc cttggggggt gtaaaagaga gtnacaggta cccccaaagg agtagatg nc 420
aaggggccct aagttgcaaa atgatgtcga ttttggggcc gggg 464

<210> 367
<211> 1476
<212> DNA/RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(1476)
<223> apr-1 protein (APR-1) gene.

<400> 367
ctggaagaat tcgcgtggca ggagaggcgg ggccaatttt gctgagcttt ctgcggggct 60
tgcagctgcg gcaagtgcgt gcggcggtct ctgcgcgaag tcagctggcg tgggaactac 120
cctttgtagc tgagaacggc ttgtttattg ctacaaagac totattgaca ttggtagctt 180
cagcggcagc agcttct tac ggtataaagc tgttgcttcc tgaagaggct acaagcatcc 240

234/292

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ttccctagga ctgctgtaag ctttgagcct ctagcaggag acatgcctcg gggacgaaag 300
agtcggcgcc gccgtaatgc gagagccgca gaagagaacc gcaacaatcg caaaatccag 360
gcctcagagg cctccgagac ccctatggcc gcctctgtgg tagcgagcac ccccgaaga c 420
gacctgagcg gcccgcagga agaccgagc actccagagg aggcctctac caccctgaa 480
gaagcctcga gcaactgcca agcacaaaag ccttcagtgc ccgggagcaa ttttcagggc 540
accaagaaaa gtctcctgat gtctatatta gcgctcatct tcatcatggg caacagcgcc 600
aaggaagctc tggctctggaa agtgctgggg aagttag gaa tgcagcctgg acgtcagcac 660
agcatctttg gagatccgaa gaagatcgtc acagaagagt ttgtgcgcag aggttacctg 720
atttataaac cggtgccccg tagcagtcog gtggagtatg agttcttctg ggggccccga 780
gcacacgtgg aatcgagcaa actgaaagtc atgcattttg tggcaagggt tcgtaaccga 840
tgctctaaag actggccttg taattatgac tgggattcgg acgatgatgc agaggttgag 900
gctatcctca attcaggtgc taggggttat tccgcccctt aagtagatct gaggcagacc 960
cttgggggtg taaaagagag tcacaggtac cccaaggagt agatgccagg gtcctaagtt 1020
gaaaatgatg tcgattgggg gcgggggaca ctgtatttga tatttgtgat cagtga tcat 1080
tgttcaactg cgaaatagag tgtttgcttt tgataatgga aaattgtatt cgttttaaaa 1140
ttcgttttgt tgagaataac aatatgttta aaaatataat tgaacaaatt tttttctttg 1200
tttctgtca ttgacattta gtataacagt tttgctaacg ttctaaaaatg aagtcgttcc 1260
atcataatct atgatcttgt acagcactta tagaaataag ctgttctttt gaagttgaaa 1320
taccagtaaa aatgttgaag aaggatggag gatttcttca tatctgacgt ttctgaaacc 1380
ctttgtgtct gctgttgtgt gaagattgac atttaccatg attttcctta gttacttgac 1440
gtcttgtatc tctttttatt ttcggattgc ttatca 1476

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<210> 368

<211> 436

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(436)

<223> 3' terminal sequence. ets variant gene 5
(ets-related molecule) (ETV5) gene.

<400> 368

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cgtttttttg ctttaaatac caaaactaca aaaatcagtt tataaactgt ttttccaaaa 60
caaccaccaa aacaaaacaa tcccccaaat cagggcaaaa caaaatactg tcaaaagtgt 120
taatcgccct tctcctaaaa taaaagtcac ccacactcag ccacgtgatt gggaagagaa 180
agggggcttg ctctacttgg cgaccacatg gccgggtggg tcccaa gagt agccatgggt 240
tatgatcttg agaaccacgg agngcgaaac agctgttctg actgcccccc tttttctaga 300
caaggggtaa tatttcagat tcagctagaa gagctttcca atgtttaaga tgtattttta 360
accttaatg gtttgagcct cccaactta gcctacttac ttttcaagg gtttgtgatt 420
tttcaacaaa ttgtgc 436

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<210> 369

<211> 414

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(414)

<223> 5' terminal sequence. ets variant gene 5
(ets-related molecule) (ETV5) gene.

<400> 369
 gggttgctogg cgctggggca tccagaagaa cgggccagcc atgaactatg acaagctgag 60
 ccgctctctc cgctattact atgaaaagg catcatgcag aaggtggctg gagagcgata 120
 cgtctacaaa tttgtctgtg acccagatgc cctcttctcc atggctttcc cggataacca 180
 gcgtccgttc ctgaaggcag agtccgagtg ccacctcagc gaggaggaca cctgccoget 240
 gaccactttt gaagacagcc ccgcttacct cctggacatg gaccgctgca gcagcctccc 300
 ctatgccgaa ggtttgctta ctaagtttct gagtggcgga gtgnccaaac cctaggagct 360
 agcagttccc attcagggca aacaagnggc agtgnggttt gtt ttgtgtt tttt 414

<210> 370
 <211> 249
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(249)
 <223> 5' terminal sequence. cd69 antigen (p60,
 early t-cell activation antigen) (CD69) gene.

<400> 370
 ataataagga aacgtgttca cttattgact attatagaat ggaactcatg gaaatctgtg 60
 tcagtggatg ctgctctgtg gtccgaagtc ttccatagag actttgtgaa aaaaaatttt 120
 atagtgtctt gggaattttc ttccaaacag aactatggaa aaaaaggaag aaattccagg 180
 aaaatctgca ctgtgggctt ttattgccat gagctagaag catcacaggg tgaccaataa 240
 cccngacgc 249

<210> 371
 <211> 1702
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(1702)
 <223> cd69 antigen (p60, early t-cell activation
 antigen) (CD69) gene.

<400> 371
 agactcaaca agagctccag caaagacttt cactgtagct tgacttgacc tgagattaac 60
 tagggaatct tgagaataaa gatgagctct gaaaattggt tcgtagcaga gaacagct ct 120
 ttgcatccgg agagtggaca agaaaatgat gccaccagtc ccattttctc aacacgtcat 180
 gaagggtcct tccaagttcc tgtcctgtgt gctgtaatga atgtggtctt catcaccatt 240
 ttaatcatag ctctcattgc cttatcagtg ggccaatata attgtccagg ccaatacaca 300
 ttctcaatgc catcagacag ccattgtttc tcatgc tctg aggactgggt tggctaccag 360
 aggaaatgct actttatttc tactgtgaag aggagctgga cttcagccca aaatgcttgt 420
 tctgaacatg gtgctactct tgctgtcatt gattctgaaa aggacatgaa ctttctaaaa 480
 cgatacgag gtagagagga acactgggtt ggactgaaaa aggaacctgg tcacctatgg 540
 aagtgggtcaa atggcaaaga atttaacaac tgggtcaacg ttacagggtc tgacaagtgt 600
 gttttctga aaaacacaga ggtcagcagc atgggaatgtg agaagaattt atactggata 660
 tgtaacaaac cttacaaata ataaggaaac atgttcactt attgactatt atagaatgga 720
 actcaaggaa atctgtgtca gtggatgctg ctctgtgggc cgaagtcttc cataga gact 780

236/292

```

ttgtgaaaaa aaattttata gtgtcttggg aattttcttc caaacagaac tatggaaaaa 840
aaggaagaaa ttccaggaaa atctgcactg tgggctttta ttgccatgag ctagaagcat 900
cacagggttg ccaataacca tgcccaagaa tgagaagaat gactatgcaa cctttggatg 960
cactttatat tattttgaat ccagaaataa tgaa ataact aggcgtggac ttactattta 1020
ttgtgtaatg actaccaaca gtgagagccc ttcattgcatt tgcactactg gaaggagtta 1080
gatgttggta ctagatactg aatgtaaaca aaggaattat ggctggtaac atagggtttt 1140
agtctaattg aatcccttaa actcaggagg cattttataaa tggacaaatg cttatgaaac 1200
taagatttgt aatattttct tctttttaga gaaatttgcc aatttacttt gttatttttc 1260
cccaaaaaga atgggatgat cgtgtattta tttttttact tcctcagctg tagacaggtc 1320
cttttcgatg gtacatatat ctttgccctt ataatctttt atacagtgtc ttacagagaa 1380
aagacataag caaagactat gaggaatatt tgcaagacat agaatagt gt tggaaaatgt 1440
gcaatatgtg atgtggcaaa tctctattag gaaatattct gtaatottca gacctagaat 1500
aatactagtc ttataatagg tttgtgactt tcctaaatca attctattac gtgcaatact 1560
tcaatacttc atttaaaata tttttatgtg caataaaatg tatttgtttg tattttgtgt 1620
tcagtacaat tataagctgt tt ttatatat gtgaaataaa agtagaataa acacaaaaaa 1680
aaaaaaaaaa aaaaaaaaaa aa

```

1702

<210> 372

<211> 585

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(585)

<223> 3' terminal sequence. oncogene tc21 (TC21) gene.

<400> 372

```

gtaggcagta tgattccaaa agttaaaaat tatttcacaa cctgtagctt cagcttggca 60
aacagcttag attccaaaac tgattcatct ctattaaaat gtaagcactt aaaaaaagag 120
catgtctgtg tatatagaca tat attttaa aggaatcaga taatctttga agcagcctta 180
gtgtttcctt taaatttgte tggaaatgac cattgtatta gottcacaga aaggactagc 240
cagcttcttc gtctaaggct aacatggtga tcatttgtct aaggctagaa aggtaccaac 300
aagatgtaaa ctgaggagag aaagagaaga tgagggcttt tcctggccgt tggtagctaa 360
aactgaaggg attctagaaa atgacacaat ggcagccttt ontgtctttt tctttccgtg 420
ttgggttcngg tgaaggagga cattcctggc cctggaaatt tccnggataa ccgggacaag 480
ttcatgggaa agcttgatct acattcatcc taatccttgc cggatgccnc catgtatgtt 540
acctaagctg ccggcaacgg tngcctctnc cggggtaccg gcc ng

```

585

<210> 373

<211> 451

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(451)

<223> 5' terminal sequence. oncogene tc21 (TC21) gene.

<400> 373

```

gattcttaca caaagcagtg tgtgatagat gacagagcag ccgggctaga tattttggat 60

```


237/292

```

acagcaggac aagaagagtt tggagccatg agagaacagt atatgaggac tggcgaaggc 120
ttcctgtttg tcttttcagt cacagataga ggcagttttg aagaaatcta taagtttcaa 180
agacagattc tcagagtaaa ggatcgtgat gagttcccaa tgattt taat tggtaataaa 240
gcagatctgg atcatcaaaag acaggtaaca caggaagaag gacaacagtt agcacggcag 300
cttaaggtaa catacatgga ggcacagca aagattagga tgaatgtaga tcaantttcc 360
atgaacttgt ccgggttatc aggaaatttc aagancagga atgtcctctt cacagaccac 420
acggaaagaa aagccagaaa gctg cattgt g 451

```

<210> 374

<211> 425

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(425)

<223> 3' terminal sequence. cd44 antigen (homing
function and indian blood group system) (CD44)
gene.

<400> 374

```

gaagatcgaa gaagtacaga tatttattat gaatcagttt aaaccctttt gtgcctctga 60
caaagtaact ttaaaaaaatt atactgatca aaggactgat ccagggttta atatttcaaa 120
aacacagata aatagttttac tacagataaa tagcttcacc c tttggtgtc ctcccagaag 180
catctgaaaa atttctagag ggggtctgtt gaagatgtgt aactagtaca ccccaacccc 240
caacctcagt ggaaagcaat gccagggat taggctatgg aagggcacaaa tggaccatt 300
caaatttcct ccaggggacc aggcctatt aaccctggga aatgtcctta gctggtgggg 360
gaaaggttgg cgattcagga atacatatgt gtagtttttg ttagaagcca tccatagcac 420
acccg 425

```

<210> 375

<211> 478

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(478)

<223> 5' terminal sequence. cd44 antigen (homing
function and indian blood group system) (CD44)
gene.

<400> 375

```

ggcgttccag ttcccacttg gaggccttc atccctcggg tgtgctatgg atggcttcta 60
acaaaaacta cacatatgta ttcctgatcg ccaacct ttc cccaccagc taaggacatt 120
tcccagggtt aatagggcct ggtccctggg aggaaatttg aatgggtcca ttttgccctt 180
ccatagccta atccctgggc attgctttcc actgaggttg ggggttggg tgtactagt 240
acacatcttc aacagacccc ctctangaaa tttttcagat gcttctggga gacacccaaa 300
ggggaaagct atttatctgt agtaaacat tttctgtgt ttttgaaata ttaaaccctg 360
gatcagtoct ttgatcagta taaattttt aaagttactt ttgtcagagg caccaaaggg 420
tttaaaactga ttcataaata aatatcngga cttcctcgat cttccaaaaa aaaaaaaa 478

```

<210> 376
 <211> 1794
 <212> DNA/RNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1794)

<223> cd44 antigen (homing function and indian blood group system) (CD44) gene.

<400> 376

```

ccccgcgcct ccgttcgctc cggacacccat ggacaagttt tgggtggcag ca gcctgggg 60
actctgcctc gtgccgctga gcctggcgca gatcgatttg aatataacct gccgctttgc 120
aggtgtatcc cagctggaga aaaatggctg ctacagcacc tctcggacgg aggcgcctga 180
cctctgcaag gctttcaata gcaccttgcc cacaatggcc cagatggaga aagctctgag 240
catcggattt gagacctgca ggtatgggtt c atagaaggg catgtgggtga ttccccggat 300
ccaccccaac tccatctgtg cagcaaacaa cacaggggtg tacatcctca catacaaac 360
ctcccagtat gacacatatt gcttcaatgc ttcagctcca cctgaagaag attgtacatc 420
agtcacagac ctgcccattg cctttgatgg accaattacc ataactattg ttaaccgtga 480
tggcaccggc tatgtccaga aaggagaata cagaacgaat cctgaagaca tctaccccag 540
caaccctact gatgatgacg tgagcagcgg ctccctccagt gaaaggagca gcacttcagg 600
aggttacatc ttttacacct tttctactgt acaccccatc ccagacgaag acagtccctg 660
gatcacccgac agcacagaca gaatccctgc taccagagac caagacacat t ccaccccag 720
tgggggggtcc cataccactc atggatctga atcagatgga cactcacatg ggagtcaaga 780
aggtggagca aacacaacct ctggctctat aaggacaccc caaattccag aatggctgat 840
catcttggca tccctcttgg ccttggcttt gattcttgca gtttgattg cagtcaacag 900
tcgaagaagg tgtgggcaga agaaaaagct agtgatcaac agtggcaatg gagctgtgga 960
ggacagaaag ccaagtggac tcaacggaga ggccagcaag tctcaggaaa tgggtgattt 1020
ggtgaacaag gagtcgtcag aaactccaga ccagtttatg acagctgatg agacaaggaa 1080
cctgcagaat gtggacatga agattggggt gtaacaccta caccattatc ttggaaagaa 1140
acaaccggtt gaaacataac cattacaggg agctgggaca cttacagatg gcaatgtgct 1200
actgattgtt tcattgcgaa tcttttttag cataaaattt tctactcttt ttgttttttg 1260
tgttttgttc tttaaagtca ggtccaattt gtaaaaacag cattgctttg taaattaggg 1320
cccaattaat aatcagcaag aatttgatcg ttcagttcca cttg gaggcc ttcacccctg 1380
ggtgtgctat ggatggcttc taacaaaaac tacacatatg tattcctgat cgccaacctt 1440
tccccacca gctaaggaca tttcccaggg ttaatagggc ctgggtccctg ggaggaaatt 1500
tgaatgggtc cattttgccc ttccatagcc taatccctgg gcattgcttt ccaactgagg 1560
tggggtgtac tagttacac a ttttcaacag accccctcta gaaatttttc agatgcttct 1620
gggagacacc aaagggtgaa gctatttata tgtagtaaac tatttatctg tgtttttgaa 1680
atattaaacc ctggatcagt cttttgatca gtataatttt ttaaagttac tttgtcagag 1740
gcacaaaagg gtttaaactg attcataata aatatctgta cttcttcgat cttc 1794

```

<210> 377
 <211> 452
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(452)

<223> 3' terminal sequence. cyclin -dependent kinase inhibitor 3 (cdk2 -associated dual specificity phosphatase) (CDKN3) gene.

<400> 377

```

ttttgtcaat aaaacttttag gaatatctgc acatgtacat ttacattcaa gttgataaca 60
ctgggtggttt catttcaata caaattatgc tagagaactg acatttcaga catgggcata 120
tatatgctat ttgaattcct ttatcttoga tacagatctt gattgtgaat ctc ttgatga 180
tagatgtgca gctaatttgt cccgaaactc atgaagataa ttgtattgct tgatggctctg 240
tattgccccg gatactctta ggtctgcag gctgtctatg gcttgcctctg gtgatattgt 300
gtcagacagg tatagtagga gacaagcagc tacaagacaa gatctcccaa gtctccata 360
gcagtgtatt aagggttttc cggtaatttt t aaggcagggt tgtaagcnet tccattattt 420
cacagcagct ggccatgten ggagtcccc ca 452

```

<210> 378

<211> 472

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(472)

<223> 5' terminal sequence. cyclin -dependent
kinase inhibitor 3 (cdk2 -associated dual
specificity phosphatase) (CDKN3) gene.

<400> 378

```

ggcacgagcg gcaactggtc tcgacgtggg ggggccanga ctgaagccca ngnttcaata 60
caaaacaagt agtttgactc atcagatg aa gagcctattg aagatgaaca gactccaatt 120
catatatcat ggctatcttt gtcacgagtg aattgtcttc agtttctcgg tttatgtgct 180
cttccaggtt gtaaaatttaa agatgttaga agaaatgtcc aaaaagatac agaagaacta 240
aagagctgtg gtatacaacg acatatitgt tttctgcacc agaggggaac tgtcaaaata 300
tagagtccca aaccttcttg atctctacca gcaatgtgga attatcacc atcatcatcc 360
aatccgcaga tggagggact cctgacatag ccagctgctg tgaaataatg gaagagctta 420
caacctgcct taaaaattac cgaaaaaact taataactg ctatggagga ct 472

```

<210> 379

<211> 639

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(639)

<223> cyclin-dependent kinase inhibitor 3
(cdk2-associated dual specificity phosphatase)
(CDKN3) gene.

<400> 379

```

atggagccgc ccagttcaat acaaacaagt gagtttgact catcagatga agagcctatt 60
gaagatgaac agactccaat tcatatatca tggctatctt tgtcacgagt gaattgttct 120
cagtttctcg gtttatgtgc tcttccaggt tgtaaattha aagatgttag aagaaatgtc 180
caaaaagata cagaagaact aaagagctgt ggtatacaag acatatitgt tttctgcacc 240
agaggggaac tgtcaaaata tagagtccca aaccttcttg atctctacca gcaatgtgga 300
attatcacc atcatcatcc aatcgcagat ggagggactc ctgacatagc cagctgctgt 360
gaaataatgg aagagcttac aacctgcctt aaaaattacc gaaaaacctt aataactg 420

```

240/292

tatggaggac ttgggagatc ttgtcttcta gctgcttgt c tctactata cctgtctgac 480
 acaatatcac cagagcaagc catagacagc ctgcgagacc taagaggatc cggggcaata 540
 cagaccatca agcaatacaa ttatcttcat gagtttcggg acaaattagc tgcacatcta 600
 tcatcaagag attcacaatc aagatctgta tcaagataa 639

<210> 380

<211> 487

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(487)

<223> 5' terminal sequence. max-interacting
protein 1 (MXI1) gene.

<400> 380

aagtggcgac tggaaacagct gcagggtcct caggagatgg aacgaatacg aatggacaga 60
 attggatcaa ctatttcttc agatcgttct gattcagagc gagaggagat tgaagtggat 120
 gttgaaagca cagagttctc ccatggagaa gtggacaata taagtaccac cagcatcagt 180
 gacattgatg accacagcag cctgccgagt attgggagtg acgagggtac tccagtgcc 240
 gtgtcaaaact ttcattcact tcatagaacc cagcatgaca taacagtgc gggaaaatat 300
 tcaactgggcc attcatataa acaatctott aaattgggtt catgatgcag tctctctttt 360
 aaaacaaaac aaaacaaaac aaactatact tgaacaaaag ggtcagagga ctgttttaag 420
 caatacttag caaaagtggg cagctcccaa gagacaattt cagatttcat ttggaaatcc 480
 catttta 487

<210> 381

<211> 2416

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2416)

<223> max-interacting protein 1 (MXI1) gene.

<400> 381

agattatgat cgctgaggc ccctctcta cccagatacc gatgttatac tgatgtgttt 60
 ttcctttttt tttttttttt tttaagtaat taagggtagt taaattattt aaagtataca 120
 aagtccaaac agccaggggt aaggtctcca agaggccttc ccagggttaag ggagtgcgga 180
 gaggcccccg tcgccaccg cgggtcccat ggagcgggtg aagatgatca acgtgcagcg 240
 tctgctggag gctgccgagt ttttgagcgc ccgggagcga gagtgtgaac atggctacgc 300
 ctcttcattc cgtcccatgc cgagcccccg actgcagcat tcaaagcccc caggaggtt 360
 gagccgggca cagaaacaca gcagcgggac gagcaacacc a gcactgccca acagatctac 420
 acacaatgag ctggaaaaga atcgacgagc tcatctgcgc ctttgttttag aacgcttaaa 480
 agttctgatt ccactaggac cagactgcac ccggcacaca acacttggtt tgctcaacaa 540
 agccaaagca cacatcaaga aacttgaaga agctgaaaga aaaagccagc accagctcga 600
 gaatttgga cgagaacaga gatttttaaa gtggcgactg gaacagctgc agggctctca 660
 ggagatggaa cgaatacgaa tggacagcat tggatcaact atttcttcag atcgttctga 720
 ttcagagcga gaggagattg aagtggatgt tgaaagcaca gagttctccc atggagaagt 780
 ggacaatata agtaccacca gcacagtgat cattgatgac cacagcagcc tgccgagtat 840
 tgggagtgac gagggttact ccagtgccag tgtcaaaact tcattcactt catagaaccc 900

241/292

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agcatgacat aacagtgcag ggcaaaatat tcactgggcc aattcaatac aaacaatctc 960
ttaaattggg ttcattgatgc agtctcctct ttaaaacaaa acaaaacaaa aaaaaactat 1020
acttgaacaa aagggtcaga ggacctgtat ttaagcaaa t acttagcaaa aagtggggca 1080
gagctcccaa ggagaacaaa tattcagaat attcatattg gaaaaatcac aatttttaat 1140
ggcagcagaa aacttggtgt aaattttctt gatttgagtt gattgagaag aggacattgg 1200
agatgccatc ctctttctct tttctcgttt gtcatacta cattgagtag acacatttaa 1260
ggatgggggt atgaaccctt cctgagcttt atggctctaa aagcaaaata aaaactattc 1320
gaatgaaaag acaagaaaat caggatttaa tcttgatag ctaataatga gctattaaaa 1380
ctcagcctgg gacagtttat catgaagcct gtggatgatc aatcctttat tattattttt 1440
tttttttgaa aaaagctcat ttcattgctc gcaaaaggag agactcccat ga agcctttt 1500
gaaagggatc atcatgcagc tcaactttct gttggattcc atgctaagca agctaacctt 1560
atcctgcatt gtttagcata ggcaaccagc tgccacctct ccactctgct gcccttaggc 1620
cacatgggag cagtccatgc atgacagcct ctatcctaca aggcctatga gtatggattg 1680
ggggggccaa aaggaaaaag ctccatg tgc ctctttgtct gcgtgggtca gaagagtgt 1740
gcacgcagat tagcaggcca aggtctgagc cacagcagca tttttatttc agattttgat 1800
aactgtttat atgtgttgaa aacccaaatg acatctttt aaagcttctc cataaaaaaa 1860
aatagatgtc ttttatagtg gaaaaacaca tggggaaaaa aatcatctat tttgatgcag 1920
catttgataa tgataaaaca cctcacacct cactctttat agtgacaaaa atgaatgagg 1980
tctgggctag gtagaaaaag ggtcaatgct atttttgttt ttagaatcat taccttttac 2040
cagcttttaa ccactctgata tctatagtag acacactatc atagttaaca tagttaagtt 2100
cagcacttgt ctcattttaa tgtaaagatt tgcttccatt ttcctacagg cagtctctct 2160
cttctcaca gtcccactgt gcagggtgcta ttgttactct tacgaatatt ttcagtaagt 2220
ttattttctt ctaagtgaag tttctagcct gcactttgat gtcattgtgt ccctttgtct 2280
ttcaaacctc aaggttcccc tgtggccctc tcccttacc tggaaggcc tcttgagac 2340
cttacccttg gctgtttgga ctttgtatc tttaaataat ttaactacc ttaattactt 2400
aaaaaaaaa aaaaaa 2416

```

<210> 382

<211> 378

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(378)

<223> 3' terminal sequence. homeo box a5 (HOXA5) gene.

<400> 382

```

tttttttttt ttgttatagt tacttcaagt aacacagctt gcttcatata aataagttaa 60
aacatctatt ttttttcaag acaaagccat tcaggacaaa gagatgaaca gaaagcagat 120
ctacttatac aggcgc tata atggcaataa acaggetcat gattaaaaga tgaattaggg 180
caacgagaac agggcttctt cacagaagga acacaaggga gtttcagaaa gtcaccttag 240
tactgacact acgcgggatc cgctaaatac tgctcagtag tttaaagcgt cagatactca 300
gggacggaag gccctccctt gcccgcggnc atnctccatg gcttttcagc ttattatc tt 360
ttttccactt caatcncc 378

```

<210> 383

<211> 439

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

242/292

<221> misc_feature
 <222> (1)..(439)
 <223> 5' terminal sequence. homeo box a5 (HOXA5) gene.

<400> 383
 aaatcaagca cacatantan aaaacaaatg agctcttatt ttgtaaactc attttgcggt 60
 cgctatccaa atggcccga ctaccagttg cataattatg gagatcatag ttccgtganc 120
 gagcaattca gggactcggc gagcatgcac tccggcaggt acggctacgg ctacaatggc 180
 atggatctca gcgtcggcng ctcgngctcc ngcacttttg ctccggagag cgcgcccga 240
 gctacgtnc aagcccaggt acagcnagcc ggccacgtcc 300
 acgcactctn cctcancncg atccgctgcn ctgctccgnc gtngggccct tcgccnnga 360
 ancgacanna ccaanggcgg gaaaaactcc cttaaggca a ctccagcngg cgctcgggc 420
 cgacngccgg aagcaccca 439

<210> 384
 <211> 813
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(813)
 <223> homeo box a5 (HOXA5) gene.

<400> 384
 atgagctctt attttgtaaa ctcatcttgc ggtcgctatc caaatggccc ggactaccag 60
 ttgcataatt atggagatca cagttccgtg agcgagcaat tcagggactc ggcgagcatg 120
 cactccggca ggtacggcta cggctacaat ggcattggatc tcagcgtcgg ccgctcgggc 180
 tccggccact ttggctcgg agagcgcgcc cgcagctacg ctgccagcgc cagcgcggcg 240
 cccgcggagc ccaggtacag ccagccggcc acgtccacgc actctcctca gcccgatccg 300
 ctgccctgct ccgcctgggc cccctcgccc ggcagcgaca cgcaccacgg cgggaaaaac 360
 tccctaagca actccagcgg cgctcggcc gacgcggca gc accacat cagcagcaga 420
 gagggggttg gcacggcgtc cggagccgag gaggacgccc ctgccagcag cgagcaggcg 480
 agtgcgcaga gcgagccgag cccggcgccg cccgcccaac ccagatcta cccctggatg 540
 cgcaagctgc acataagtca tgacaacata ggcggcccgg aaggcaaaag ggcccgagc 600
 gcctacacgc gctaccagac cctggagctg gagaaggagt tccacttcaa ccgttacctg 660
 accgcagaa ggaggattga aatagcacat gctctttgcc tctccgagag acaaattaaa 720
 atctggttcc aaaaccggag aatgaagtgg aaaaaagata ataagctgaa aagcatgagc 780
 atggccgcgg caggaggggc cttccgtccc tga 81 3

<210> 385
 <211> 447
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(447)
 <223> 3' terminal sequence. x-box binding protein 1 (XBP1) gene.

<400> 385

243/292

```

gcattgtacc ttttaattgc atgggtagtt ttaaataaat ggagaaagca cttttcagaa 60
gctacactag caggaaaaaa ttccatcaag catttacata gtaaatttct ataatttcac 120
aaaagattct tgatcttact tgaagtatac atgagggaaa gagccccctc agcaggtggt 180
cccgtttgctt acagaagcaa actaaaggac ctaaaactgg aggcaagcca ggatgc caaa 240
aagggggaag agaaatgata aagaaccatt cataaattcc atgtctactt caagacattt 300
gtctaattgac cttacataa taagtatttt agggaaaact accacccttt taagataaaa 360
gtacaatctt aaaagctgta gttctcaatt atagtaatat ttontacttc cagtaatatg 420
tctcaatacc ttggactgct ggatgctc                                     447

```

<210> 386

<211> 462

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(462)

<223> 5' terminal sequence. x-box binding protein

1 (XBP1) gene.

<400> 386

```

aagaacctgt agaagatgac ctctgtccgg agctgggtat ctcaaactctg ctttcatcca 60
gccactgccc aaagccatct tctgcctac tggatgctta cagtgactgt ggatacgggg 120
gttccctttc cccattcagt gacatgtcct ctctgcttgg tgtaaaccat tcttgggagg 180
acacttttgc caatgaactc tttcccca gc tgattagtgt ctaaggaatg atccaatact 240
gttgcctttt tctttgacta ttacactgcc tggaggatag cagagaagcc tgtctgtact 300
tcattcaaaa agccaaaata gagagtatac agtcctagag aattcctcta ttgttcaga 360
tctcatagat gacccccagg tattgtcttt tgacatccca agcagtcocaa ggtattggag 420
acataattact gggaagtaaa gaaatattac tnataattgg ag                                     462

```

<210> 387

<211> 1836

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1836)

<223> x-box binding protein 1 (XBP1) gene.

<400> 387

```

ggcgctgggc ggctgcggcg cgcggtgcgc ggtgcgtagt ctggagctat ggtgggtggtg 60
gcagccgcgc cgaacccggc cgacgggacc cctaaagtct tgcttctgtc ggggcagccc 120
gcctccgcgc cggagcccc ggccggccag gccctgcgc tcattggtgcc agcccagaga 180
ggggccagcc cggaggcagc gagcgggggg ctgccccagg cgcgcaagcg acagcgcctc 240
acgcacctga gccccgagga gaaggcgctg aggaggaaac tgaaaaacag agtagcagct 300
cagactgcc a gagatcgaaa gaaggctcga atgagtgagc tggaacagca agtggttagat 360
ttagaagaag agaaccaaaa acttttgcta gaaaatcagc ttttacgaga gaaaactcat 42 0
ggcctttag tag ttgagaacca ggagttaaga cagcgcttgg ggatggatgc cctggttgc 480
gaagaggagg cggaagccaa ggggaatgaa gtgaggccag tggccgggtc tgctgagtcc 540
gcagcactca gactacgtgc acctctgcag caggtgcagg cccagttgtc acccctccag 600
aacatctccc catggattct ggcggtattg actcttcaga ttcagagtct gatatcctgt 660
tgggcattct ggacaacttg gacccagtca tgttcttcaa atgcccttcc ccagagcctg 720

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244/292

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ccagcctgga ggagctccca gaggtctacc cagaaggacc cagttcctta ccagcctccc 780
tttctctgtc agtggggacg tcatcagcca agctggaagc cattaatgaa ctaattcgtt 840
ttgaccacat atataccaag cccctagtct tagagatacc ctctgagaca gagagccaag 900
ctaagtgtgt agtgaaaatc gaggaagcac ctctcagccc ctccagagaat gatcacccctg 960
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tctaaggaat gatccaatac tgttgccctt ttcttgact attacactgc ctggaggata 1260
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gtaagatcaa gaatcttttg tgaaattata gaaatttact atgtaaat gc ttgatgggaat 1740
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caattaaaag gtacaatgca aaaaaaaaaa aaaaaa 1836

```

<210> 388

<211> 433

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(433)

<223> 3' terminal sequence. tumor necrosis factor,
alpha-induced protein 3 (TNFAIP3) gene.

<400> 388

```

tttttcttaa ataatttatt ttttaatgtt gactcttgtg aaaagttaca tttattttaga 60
aaaacttgaa gaaatcoa ac aaagaatagg tggctttcta ttagggacaa ttaaatgtgc 120
aaatttcaaa tactttttat aataagtata aataattact ttttttcaca ttaagaatgg 180
aaataatgat caacacaana tattaagata tcaactttaa gagaattaga tgaaaacatg 240
gaagtttatt tgngtatcct tggaaangaa ttataaagat tcgtctggga aaaactt agg 300
gggctctaag gggaaagttg tgcctaatag tatgagtaaa ggctgtgtag agttatggat 360
cacaaatatt ttcaggccnt aagtacagac cccnnaaatg gcagccttta tcnccgggga 420
aatgcatatt ccc 433

```

<210> 389

<211> 206

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(206)

<223> 5' terminal sequence. tumor necrosis factor,
alpha-induced protein 3 (TNFAIP3) gene.

<400> 389

245/292

ctcaaccagc tgccttttta aaggga gctc tagtcctttt tgtgtaatto actttattta 60
 ttttattaca aacttcaaga ttatttaagt gaagatattt cttcagctct ggggaaaatg 120
 ccacagtgtt ctcttgagag aacatccttg ctttgagtca ggctgtgggc aagtctctga 180
 ccacagggag taaattngnn cctctt 206

<210> 390

<211> 4426

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(4426)

<223> tumor necrosis factor, alpha -induced protein
 3 (TNFAIP3) gene.

<400> 390

tgccttgacc aggacttggg acttttgcga aggatcgcg ggcccggaga ggtgttggag 60
 agcacaatgg ctgaacaagt ccttctcag gctttgtatt tgagcaatat gcggaaagct 120
 gtgaagatac gggagagaac tccagaagac atttttaaac ctactaatgg gatcattcat 180
 ctttttaaaa ccatgcaccg atacacactg gaaatgttca gaacttgcca gtt ttgtcct 240
 cagtttcggg agatcatcca caaagccctc atcgacagaa acatccaggc caccctggaa 300
 agccagaaga aactcaactg gtgtcgagaa gtccggaagc ttgtggcgct gaaaacgaac 360
 ggtgacggca attgcctcat gcatgccact tctcagtaca tgtggggcgt tcaggacaca 420
 gacttggtac tgaggaaggg gctgttcagc a cgctcaagg aaacagacac acgcaacttt 480
 aaattccgct ggcaactgga gtctctcaaa tctcaggaat ttgttgaaac ggggctttgc 540
 tatgatactc ggaactggaa tgatgaatgg gacaatctta tcaaatggc ttccacagac 600
 acacccatgg ccgaagtgg acttcagtac aactcactgg aagaaatata catatttgtc 660
 ctttgcaaca tctcagaag gccaatcatt gtcatttcag acaaatgct aagaagtgtg 720
 gaatcagggt ccaatttcgc ccttttgaaa gtgggtggaa tttacttgcc tctccactgg 780
 cctgcccagg aatgctacag ataccccat gttctcggct atgacagcca tcattttgta 840
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 gacgggggaa gatttgaaga cttaaaagt cactttttga cagatcctga aaatgagatg 960
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 cccaaccaga ggatgggccc tggggccccc cggggtgagc ctgcccccca agaccccccc 2340
 aagcagcggt gccgggcccc cgccctgtgat ctttttgga atgccaagtg caacggctac 2400

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tgcaacgaat gcttttcagtt caagcagatg tatggctaac cggaaacagg tgggtcacct 2460
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```

<210> 391

<211> 440

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(440)

<223> 3' terminal sequence. serum response factor
(c-fos serum response element-binding
transcription factor) (SRF) gene.

<400> 391

```

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agttagaaac actgacatac aactcacat tcaagcacac aactcactc aggcgcacac 180
accacacac acatacccca gagccaccga ggaagggaaa caccaagggt cgctgcacat 240
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aacacacatt ataagcactt tgctgatto actcactnng gtctgtcttt tgtgggaagg 360
agaggaagaa ttcattcaaag gtctcctccc catgggtngg gggagtgggg agtgagttag 420
tgatggtgga gtgaaacaag

```

440

<210> 392
 <211> 471
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(471)
 <223> 5' terminal sequence. serum response factor
 (c-fos serum response element-binding
 transcription factor) (SRF) gene.

<400> 392
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 tgcagcaaag gggagccaga aatgggcagt tctcccaggg agtgagcagc tactgttaact 120
 ttttttaaatt aagacaaaaa gccttgaaga aaatgacttt atttttctaa gtgtaacctc 180
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 ccttgagggt gggccagcat aggggggagg gtcttttacc ctgtgtcaga gctaccttc 300
 accacctata tccagaaggg gagctttttc agaaacaggg cagcagtggg gtgaaatatt 360
 cttaacccct aagactgcct tcagtaagga acaagctggc ttctgtgatt aggtgaaggg 420
 atgggggaag attttaatgc acagccta gt.tatcaagggg atgatttgcc g 471

<210> 393
 <211> 4201
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(4201)
 <223> serum response factor (c-fos serum response
 element-binding transcription factor) (SRF) gene.

<400> 393
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 ccgcgccag cagccctgc cccccgggg acgctgacgg ccgcccggcg cgcgccta 120
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 gtgagggccc aggtcggccc gggcgtgcag gggccccggg ttgcagcgg cggccgccc 240
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 gccagactct ccaccccggt cag accccac aacagaccag agaatgagtg ccaactggctt 1080
 tgaagagaca gatctcacct accaggtgtc ggagtctgac agcagtgggg agaccaagga 1140

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cacactgaag cgggcgttca cagtcaccaa cctgccgggt acaacctcca ccatccaaac 1200
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agaagtggga gcagcttctt gggctgagt cagccaaagg ggagccagaa atgggcagtt 3180
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cccaccccc tctccccct ttggggtaga ccttgagggt gggccagcat aggggggagg 3360
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g 4201

```

<210> 394

<211> 563

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(563)
 <223> 3' terminal sequence. sry (sex determining region y)-box 9 (campomelic dysplasia, autosomal sex-reversal) (SOX9) gene.

<400> 394
 tttttaatgc aatgtatatt tattgtaaac aataatatac aaaaaaaaaa aagagaaaga 60
 aaaagggaaa ggtaagtttc acggagagaa caaaagggtt ggggctggga gggaacaag 120
 tgaacaaac aaaacacgaa cacaaccaa agcttttacc taaagacaaa atatgattta 180
 aatgccaggt ttcttaagtt acaga agtat ctttttaaaa agatctgctt ttatacagaa 240
 attgaaggat gccatattat gagtgcttta agattttatt ctactgactt ctaaaactgt 300
 taatataatct ttttttaaat aaaaaaaaaa gtttgctgtc ttttttaaaa agcaatcctc 360
 aaactctcta gccacagcag taattaagat taaggctctgt cagtgggctg atccccctca 420
 ggtagcctcc ctactccaa gagaagatgc ngagaaatat gggatggaca catgcctgca 480
 tgtttttgtg nccaaacaca cacanacca nacacacnca caatataagg cngccccaag 540
 gtcntggcc gaaancctgg caa 563

<210> 395
 <211> 3936
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(3936)
 <223> sry (sex determining region y) -box 9
 (campomelic dysplasia, autosomal sex -reversal)
 (SOX9) gene.

<400> 395
 ggagagccga aagcggagct cgaaactgac tggaaacttc agtggcgcgg agactcgcca 60
 gtttcaaccc cggaaacttt tctttgcagg aggagaagag aaggggtgca agcgcccca 120
 cttttgctct ttttctctcc ctctctctcc tctccaattc gcctccccc acttgagcg 180
 ggcagctgtg aactggccac ccgcgcctt cctaagtgtc cgccgcggta gccggc cgac 240
 ggcgcagctt ccccgggagc cgcttgctcc gcctccgggc agccgagggg agaggagccc 300
 gcgcctcgag tcccgagcc gcgcgcgctt ctgcctttc ccggccacca gccccctgcc 360
 ccgggcccgc gtatgaatct cctggacccc ttcataaga tgaccgacga gcaggagaa 420
 ggctgtccg gcgccccag cccaccatg tccg aggaact ccgcgggctc gccctgccc 480
 tggggtccg gctcggacac cgagaacacg cggccccagg agaacacgtt cccaagggc 540
 gagcccgatc tgaagaagga gagcgaggag gacaagttcc ccgtgtgcat ccgcgagcg 600
 gtcagccagg tgctcaaagg ctacgactgg acgctgggtc ccatgccggt gcgcgtcaac 660
 ggctccagca agaacaagcc gcacgtcaag cggcccatga acgccttcat ggtgtggg 720
 caggcggcgc gcaggaagct cgcggaccag tacccgact tgcacaacgc cgagctcagc 780
 aagacgctgg gcaagctctg gagacttctg aacgagagcg agaagcggcc cttcgtggag 840
 gaggcggagc ggctgcgcgt gcagcacaag aaggaccacc cggattacaa gtac cagccg 900
 cggcggagga agtcggtgaa gaacgggcag gcggaggcag aggaggccac ggagcagacg 960
 cacatctccc ccaacgccat cttcaaggcg ctgcaggccg actcgccaca ctctctctcc 1020
 ggcatgagcg aggtgcactc ccccggcgag cactcggggc aatcccaggg cccaccgacc 1080
 ccaccacca ccccaaaac cgacgtgcag ccgggcaagg ctgacctgaa gcgagagggg 1140
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 gagctgagca gcgagctcat ctccaacatc gagaccttgc atgtcaacga gtttgaccag 1260
 tacctgccgc ccaacggcca cccgggggtg ccggccacgc acggccaggt cacctacag 1320
 ggagctacg gcacagcag caccggggcc acccggcgga gcgcgggcca cgtgtggatg 1380
 tccaagcagc aggcgcccgc gccacccccg cagcagcccc cacaggcccc gccggccccg 1440

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caggcgcccc cgcagccgca ggcgggcgccc ccacagcagc cggcgggcacc cccgcagcag 1500
ccacaggcgc acacgctgac cagcgtgagc agcgagccgg gccg gtccca gcgaacgcac 1560
atcaagacgg agcagctgag ccccgagccac tacagcgagc agcagcagca ctgcgcccaa 1620
cagatgcgct acagccgctt caacctccca cactacagcc cctcctaccc gcccatcacc 1680
cgctcacagt acgactacac cgaccaccag aactccagct cctactacag ccacgcgcca 1740
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tacaccccca tcgcccgcac ctctggggtc ccttccatcc cgcagaccca cagccccag 1860
cactgggaac aaccgctcta cacacagctc actcgacctt gaggaggcct cccacgaagg 1920
gcgacgatgg ccgagatgat cctaaaaata accgaagaaa gagaggacca accagaat tc 1980
cctttggaca tttgtgtttt tttgtttttt tattttgttt tgttttttct tcttcttctt 2040
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gacctatcca agcgcttac ccacttgtgg ccaatcagtg gccaggccaa ccttggtctaa 2160
atggagcagc gaaatcaacg agaaactgga ct ttttaaac cctcttcaga gcaagcggtg 2220
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gtaagcttta tgatataat attttttaa gaagagaaaa acaccttgag ccttaaaacg 2760
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tgcagtctta agaaagaggt aaaaggcaag caaaggagat gaaatctggt ctgggaatgt 2880
ttcagcagcc aataagtgc cgagcacact gcccccggtt gcctgctgg gccccatgtg 2940
gaaggcagat gctgctcgc tctgtcacct gtgcctctca gaacaccagc agttaacctt 3000
caagacattc cacttgctaa aattatttat tttgtaagga gagggtttta ttaaaacaaa 3060
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aagatacttc tgtaacttaa gaaacctggc atttaaatca tttttgtct ttaggtaaaa 3780
gctttggttt gtgttctgtt tttgtttgt tcaactgttt cctcccagc ccaaacctt 3840
ttgttctctc cgtgaaactt acctttccct ttttctttct cttttttttt ttgtatatta 3900
ttgtttacaa taaatatata ttgcattaaa aagaaa 3936

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<210> 396

<211> 204

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(204)

<223> 3' terminal sequence. cadherin 15,
m-cadherin (myotubule) (CDH15) gene.

<400> 396

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tttttttttt tttttttttt tttttttttt ttttttttca ttcagattta cccaggaggt 60
tgctgtcttt canacaaaga tgaggttcac tggaggaggg caaagggtgg actagggagg 120

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tgaccgcgat gggccagatn ggagagaaac ttttcccacc ccggcagaag gggcctcttc 180
 ctggccgccc catccanact cagg 204

<210> 397
 <211> 458
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(458)
 <223> 5' terminal sequence. cadhe rin 15,
 m-cadherin (myotubule) (CDH15) gene.

<400> 397
 caggacgcct acgacatcag ccagctgcgt caccgcagag cgctgagcct gcctctggga 60
 ccgcccgcac ttgcagaga tgccccgcag ncagcctgca ccccagcca ccccgagtgc 120
 tgcccaccag cccctggac atgcgcgact tcatcaatga tggcttggg g gctgcagata 180
 gtgaccccag tgtgcgcct tacgacacag ccctcatcta tgactacgag ggtgacggct 240
 cggtggcggg gacntgagct ccctcctgtc cagccagggc gatgaggacc aggactacga 300
 ctacctcaga gactgggggc cccgcttcgc ccggtggca gacatgtatg ggcacccgtg 360
 cggttngga gttacggggc cagatgg gac caccaggcca gggagggctt ttctcctggg 420
 gcactgctac ccagacacag aggcgggaca gcctgan 458

<210> 398
 <211> 2833
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(2833)
 <223> cadherin 15, m-cadherin (myotubule) (CDH15)
 gene.

<400> 398
 acttgcgctg tcaactcagcc tggacgcgt tcttcgggtc gggggtgcac tccggcccgg 60
 ctcccgcctc ggcccgcgat gacgcgcgt tctcctcgt cctcggtg ttggcccaga 120
 gcctctgcct gtctttgggg gtt cctggat ggaggaggcc caccacctg taccctggc 180
 gccggggccc tgccctgagc cgcgtgcgga gggcctgggt catccccccg atcagcgtat 240
 ccgagaacca caagcgtctc ccctaccccc tggttcagat caagtccgac aagcagcagc 300
 tgggcagcgt catctacagc atccagggac ccggcgtgga tgaggagccc cggggcgtct 360
 tctctatcga caagttcaca gggaaggtct tctcaatgc catgctggac cgcgagaaga 420
 ctgatcgctt caggctaaga gcgtttgccc tggacctggg aggatccacc ctggaggacc 480
 ccacggacct ggagattgta gttgtggatc agaatgacaa ccggccagcc ttcttgacgg 540
 aggcgttcac tgcccgcggt ctggagggtg cagtcccagg cac ctatgtg accagggcag 600
 aggccacaga tgccgacgac cccgagacgg acaacgcagc gctgcgggtc tccatcctgc 660
 agcagggcag ccccgagctc ttcagcatcg acgagctcac aggagagatc cgcacagtgc 720
 aagtggggct ggaccgcgag gtggtcgcgg tgtacaatct gacctgcag gtggcggaca 780
 tgtctggaga ggcctcaca g ccaactgcct cagccatcat cacccttgat gacatcaatg 840
 acaatgcccc cgagttcacc agggatgagt tcttcatgga ggccatagag gccgtcagcg 900
 gagtggatgt gggacgcctg gaagtggagg acagggacct gccaggctcc ccaaactggg 960
 tggccagggt caccatcctg gaaggcgacc ccgatgggca gttcaccatc cgcacggacc 102 0

252/292

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ccaagaccaa cgaggggtgtt ctgtccattg tgaaggccct ggactatgag agctgtgaac 1080
actacgaact caaagtgtcg gtgcagaatg agggcccgcgt gcaggcggct gcccttaggg 1140
ctgagcgggg ccaggccaag gtccgcgtgc atgtgcagga caccaacgag ccccccgtgt 1200
tccaggagaa cccacttcgg accagcctag cagagggg gc acccccaggc actctgggtg 1260
ccaccttctc tgcccgggac cctgacacag agcagctgca gaggtcagc tactccaagg 1320
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acgtgtctcag ccggcggtcc ccccttctca agggcggtg gtacagagcc atcgtcctgg 1440
cccaggatga cgctctccag cccgcaccg ccaccggcac cctgtccatc gagatcctgg 1500
aggtgaacga ccatgcacct gtgctggccc cgccgcggcc gggcagcctg tgcagcgagc 1560
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ccttcctgcc ggggtgggaa gagtttctct ccacggccc catgcgggtc acc tcctag 2760
tcccaccttt gcctctacc agtgaacctc atctttgtat gaaagacagc aacctcctgg 2820
gtaaatctga atg 2833

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<210> 399

<211> 646

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer

<220>

<221> misc_feature

<222> (1)..(646)

<223> 3' terminal sequence. b-cell cll/lymphoma 2
(BCL2) gene.

<400> 399

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tatctcacac tgtactttat ttttcttcac aatattaact agacagacaa ggaaagttaa 60
atggcaatgt gactttttcc aacaacacaa acaaagtgcc attata gcta atgggtggcca 120
actggagact tactttacct taacctatga aagtatcctt accgtatttt ttatgtgtac 180
agtgttgcag aatatcagcc acctcttaaa agtatcaatc ttaaaaagag ccatggaagg 240
taaaagtatg aaaatcttga taacaaaagc tttcaataca aaaacactta ttgtacactt 300
atTTTTatTTT aaaacaaaaa taac cccagt aactcaaaaac aaaagcaaac cttggttgaa 360
aacttaagaa ggtataataa acaaaaccac caaaagaaaag cttccccaaa agaaatgcaa 420
tccactgtca ctcttgcaaa ttctaccttg gagggaaaaa cttaatgaaa tgagctatct 480
ggagggccca cggagatttt ccaaaagggt taggtgcatg gatttactca gtatctacnt 540
acagtcttat ttattaatag ctgaganttc ctgattgagc gaggctttcc atctccacca 600
gtgtcccccac ttctgtgcnc acttgggntg cagacaccct gtgttg 646

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<210> 400
<211> 465
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(465)
<223> 5' terminal sequence. b-cell cll/lymphoma 2
(BCL2) gene.

<400> 400
ggtattttaa caataaatgt gcagttttta ctaacaggat atttaatgac aacctttctgg 60
ttggtaggga catctgtttc taaatgttta ttatgtacaa tacagaaaaa aattttataa 1 20
aattaagcaa tgtgaaactg aattggagag tgataatata agtccttttag tcttaccag 180
tgaatcattc tgttccatgt ctttgacaa ccatgacctt ggacaatcat gaaatatgca 240
tctcactgga tgcaaagaaa atcagatgga gcatgaatgg tactgtaccg gtccatctgg 300
actgccccag aaaaataact tcaagcaaac atcctatcaa caacaagggtt gttctgcata 360
ccaagctgag cacagaagat gggaacactg gtggaggatg gaaaggctcg ctcaatcaag 420
aaaattctga gactattaat aaataagact gtagtgtaga tactg 465

<210> 401
<211> 419
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(419)
<223> 3' terminal sequence. ests (EST W73386)
gene.

<400> 401
gaaggtcatt cttgcgatgg gtttattgca ggagatgatg gaccaaattg ctctgacaca 60
tgacacagct cctgggcaag cctgctgcgn gtnogcttcc catga ccccc agggccctct 120
atgcctcccc cccagggcac cctgcccact tgccccact tcatgtacca ccaagccctt 180
tccttttctg ggcaccactc ctgagcagcg tgaccagcgg cctccaagtg catgtggctc 240
agaacataaa agcatcttca acattcgtca ttgagccaaa cgaaacacag tgtttggtc 300
aagagccggc gacactngca tcc ctatcca cactggaac ctgcccttgg gcttngttga 360
ccggaggatn ggccgctctt ccttgtcatc cagcanccgg agcatgtatg tgcccaga 419

<210> 402
<211> 568
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(568)
<223> 3' terminal sequence. granzyme a (granzyme

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1, cytotoxic t-lymphocyte-associated serine
esterase 3) (GZMA) gene.

```
<400> 402
tcatgcaaat tgattttatt tgtgaaaaga ttaagaagcc acagtanatg aaaggaaacg 60
gttattttaa ctgctccctt gatagt cata attatccagt tgagggtgtt ctttgagaga 120
agaatataga caccaggccc acgaggggtc cgcatttat ttccaaggcc aaaggaagtg 180
acccctcgga aaacaccctc gcacaacaaa gggcttccag aatctccatt gcacgagtct 240
cttcacctc ggaggcttcc agcacaaccc atattcattc caatcacagg gttaaaatta 300
tagtgatttc gatcattgca gacttttctg tctatgatgg gtgatattga cttgcaactca 360
gagtatcggg acccaagatg cactattggt gagtcctgcc ccaccctggc aacttggcac 420
atgggtcctg gntttcacat caatccccct ttttagggag atgaaggata gtcacatain 480
tggtnathtt ggctttttcc ggtcagctgt aaagttttta ggtccc ctnc gcgtttggtg 540
gggcctagcc tgggagggga aanccttt 568
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<210> 403

<211> 878

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(878)

<223> granzyme a (granzyme 1, cytotoxic
t-lymphocyte-associated serine esterase 3) (GZMA)
gene.

<400> 403

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cagattttca ggttgattga tgtgggacag cagccacaat gaggaactcc tatagatttc 60
tggcatcctc tctctcagtt gtcgtttctc tcttgctaatt tctgaagat gtctgtgaaa 120
aaattatttg aggaaatgaa gtaactcctc attcaagacc ctacatggtc ctacttagtc 180
ttgacagaaa aaccatctgt gctggggctt tgattgcaa agactgggtg ttgactgcag 240
ctcactgtaa cttgaacaaa aggtcccagg tcattcttgg ggctcactca ataaccaggg 300
aagagccaac aaaacagata atgcttgta agaaagagtt tcc ctatcca tgctatgacc 360
cagccacacg cgaagggtgac cttaaacttt tacagctgac ggaaaaagca aaaattaaca 420
aatatgtgac tatccttcat ctacctaaaa agggggatga tgtgaaacca ggaaccatgt 480
gccaagttag aggggtgggg aggaactcaca atagtgcac ttgggtccgat actctgagag 540
aagtcaatat caccatcata gacagaaaag tctgcaatga tcgaaatcac tataatttta 600
accctgtgat tggaaatgaat atgggttgtg ctggaagcct ccgaggtgga agagactcgt 660
gcaatggaga ttctggaagc ccttgttgt gcgaggtgt tttccgagg gtcacttcct 720
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```

<210> 404

<211> 191

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(191)

<223> 3' terminal sequence. v-fos fbj murine

255/292

osteosarcoma viral oncogene homolog (FOS) gene.

<400> 404

gcagtgaccg tgctcctacc cagctctgct tcacagcgcc cacctgtctc cgcccctcgg 60
 cccctcgccc ggctttgcta accgccacga t gatgttctc gggcttcaac gcagactacg 120
 aggcgatcgc ctcccgtgc agcagcgctg ccccgccggg gataancctc ctttattaca 180
 attaatacnc g 191

<210> 405

<211> 245

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(245)

<223> 5' terminal sequence. v-fos fbj murine
 osteosarcoma viral oncogene homolog (FOS) gene.

<400> 405

ttttcaactt aaatgctttt attgacaatg tcttggaca ataagcaaac aatgcttaaa 60
 tttttcattc aaattcactt tccacatgtc aaaagacctc aaggtagaaa aaaataaaat 120
 aaaaatataa atatctgaga atccatctta ataaataaat taaaaacaca ataaaacggt 180
 ttcatggaaa actgttaatg tcngaacat tcagaccacc tcnacaatgn gtgntcngtn 240
 anatt 245

<210> 406

<211> 489

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(489)

<223> 3' terminal sequence. interleukin enhancer
 binding factor 1 (ILF1) gene.

<400> 406

gcgncgcgct caccgaagg ngnangtaga cagcgggtca gaggccgcct agagccggag 60
 gacaccccaa atacaaacat accacggaga gacctgggat ctgagtttca aaagggcctg 120
 tgataaaaga ctgaatcttt ttccaaatga agtagaaatg gttctgtcgt tttaaacata 180
 cacaatactt aggagacttg ttttactcag agtggaaaat tttgccaggg acaaagtcaa 240
 cacaaagaaa caaacaacaa aaaatagcca gaaagagaac agttaagtgc agctcgggtga 300
 gtcccggcag ttccttccc gcactggctc gtcctgggt tctcaagggt ccatgcggcc 360
 acagcgtccg tccacctgtc cagcggagcc acatgtgaa atgggagggt ggataaaatt 420
 catcaggcag ctgctgtaac acggaaatgt gcagatgccg gagtagcttc gtctgaactt 480
 gaacaagac 489

<210> 407

<211> 247

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(247)

<223> 5' terminal sequence. interleukin enhancer
binding factor 1 (ILF1) gene.

<400> 407

```

tgttttcagc ctatggaatg atttcctttt gtctgtcttg ttcaagtica gacgaagcta 60
ctctggcattc tgcacatttc cgtgttacag cagctgcctg atgaa tttta tccacctcca 120
tttcagcatg tggctcgctg ggacagggtg acggacgctg tggccgcatg gaaccttgag 180
aaccagggga cgagccagtg ccgggaagga actgccggga ctcaccganc tgcncctaac 240
gtttctc                                     247

```

<210> 408

<211> 3059

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(3059)

<223> interleukin enhancer binding factor 1 (ILF1)
gene.

<400> 408

```

gccccccccc cagcctcct cccctctctc cgcccgcgcg tgctcccgcc cctcgccgcc 60
gtctgctcgc tcgcccgcgc gcctccgctc ggccccctcc ctcagctccg gtgcgcggcg 120
gccgaagacc cgcggccttg gcctcggcgc gccaccggcg ccgcgcgga gcggcccggg 180
ggccctcagc caggcccatg gcggcggcgc ggccgcctct cgggcggggc accacgcggg 240
cgccggggcg ggggcgcggg gccggggggc ggtcccgccg ggccgtgggc cgtggggccg 300
ctggaggggc gcgagttcga gtatctgatg aagaagcgct cggtgaccat cggccgcaac 360
tcgtcgaggc gctcgggtga cgtgagcatg ggccactcga gcttcattct ccggcgccac 420
ctcgagatct tcacgccccg gggcggcgcg gccatggcgg ggccgctccg gagctgcgcg 480
ccgcgcagcc caggcccgac gccggcggcg acttctacct gcgctgcttg ggcaagaacg 540
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agaagagaga gaagcaggag gcgtctgagt ctccagtga a ggccgtacag ccacacatct 720
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cctcgacctc ccagccacce gtcgtgcaga cggttcacgt cgtccaccag atcccagcg 1620
tgtcggtcac cagtgtggcc ggactggccc cagcgaacac gtacactgtc tctggacaag 1680

```

257/292

```

ctgtgggtcac cccggcagcc gtgctggccc ctccaaagc agaggcccag gagaatggag 1740
accacagggga agtcaaagt aaagtagagc ctattcccgc cattg gccac gccacgctcg 1800
gcaactgccag ccgcatcatt cagacggcac agaccacccc ggtccagacg gtgaccatag 1860
tacaacagggc acctctaggt caacaccagc taccaataaa aactgtaaca caaacaggca 1920
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```

<210> 409

<211> 201

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(201)

<223> 3' terminal sequence. rho gdp dissociation
inhibitor (gdi) alpha (ARHGDI) gene.

<400> 409

```

tttttttttt tccttcaggg gcatttattt cccggtcaga aaagaagcag ggacaggcgc 60
ctctgcctga gcctggcaga cacaacacga agaccgggga tggggcgggg gaggcacagg 120
agacggctct cagcaatgtg tgcacttggc cccttggttg ttctggctg ggtcagggaa 180
ggcctgccgn ggggtggtggc a

```

201

<210> 410

<211> 297

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(297)

<223> 5' terminal sequence. rho gdp dissociation
inhibitor (gdi) alpha (ARHGDI) gene.

<400> 410

258/292

```

ggcctctgct gccctttctg tgccccccag gttctatctc ccggtcacac ccgaggcctg 60
gcttcaggag ggagcggana gccattctcc agggcccctg gttgcccctg gacgtgtgog 120
tctgtctctc cggggtggan ctggggtgtg ggatgcacgg cctcgtgggg gccggggcgt 180
cctccagccc cgtctctccc tggccagccc cc ttgtcgtg gtcgggtccc totaaccatg 240
atgccttaac atgtggagtg tacctgtggg gccctactaa gcctctaant cactgtg 297

```

<210> 411

<211> 1819

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1819)

<223> rho gdp dissociation inhibitor (gdi) alpha
(ARHGDIA) gene.

<400> 411

```

cctgaaccgc gcggccgaac cctccgggtg cccgaccag gctaagcttg agcatggctg 60
agcaggagcc cacagccgag cagctggccc agattgcagc ggagaacgag gaggatgagc 120
actcgggtcaa ctacaagccc ccggcccaga agagcatcca ggagatccag gagctggaca 180
aggacgacga gagcctgcga aagtacaagg agggcctgct gggccgcgtg gccgtttccg 240
cagaccccaa cgtccccaac gtcgtggtga ctggcctgac cctgggtgtgc agctcggccc 300
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tgtccggcat gaagtacatc cagcatatct acaggaaagg cgtcaagatt gacaagactg 480
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accaagtga cacattgtct agagccgtct cctataggtc ccccgcccca tcccgggtgt 1740
tggtgtttgt tctgccaggc tcaggcagag gcgcctgtcc ctgcttcttt tctgaccggg 1800
aaataaatgc cctgaagg 1819

```

<210> 412

<211> 306

<212> DNA

<213> Artificial Sequence

<220>

259/292

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(306)

<223> 3' terminal sequence. complement component
4a (C4A) gene.

<400> 412

```

gctgccaaaa gcctttaata tgccctgggc ccaggctgt g ttcattgaaag cggacacagc 60
agtgccttcca gcttcatggg tcccagggtt aggttcctcc cagcggagggt gggaggggcag 120
ccctcacacc tggcaccctt gagggcatac tccctggagga agtcgttgag ctgggcacag 180
gctgcccgtt ggcgggttgc tccggcacag gcgttcagag ggcattctct cgatccagct 240
attcagatcc agcaagta ct ngggggggnc cctcccaggg gcataantng gncntccag 300
anccat 306

```

<210> 413

<211> 5417

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(5417)

<223> complement component 4a (C4A) gene.

<400> 413

```

agaaggtagc agacagacag acggatctaa cctctcttgg atcctccagc catgaggctg 60
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ctcttggtct ctctctctgt ggttcactct ggg gtcccc tatcgggtgg ggtgcagctc 180
caggatgtgc cccgaggaca ggtagtgaag ggatcagtg tctgagaaa cccatctcgt 240
aataatgtcc cctgctcccc aaagggtggc ttacacctta gctcagaaa agacttcgca 300
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acgacaaaca tccagggtat caacctgtc ttctctctc gccgggggca cctcttttg 480
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gatcagaaga tgcgccgag cactgacacc atcacagtca tgggtggagaa ctc tcacggc 600
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atcccagaca tctcagagcc agggacctgg aagatctcag cccgattctc agatggcctg 720
gaatccaaca gcagcaccca gtttgagggt aagaaatatg tcttcccaa ctttgagggtg 780
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```

```

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```


<210> 414
<211> 408
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(408)
<223> 3' terminal sequence. cd3g antigen, gamma
polypeptide (tit3 complex) (CD3G) gene.

<400> 414
ttacaatttt ccccaatagg tggcgcttct gaaacacagt atttgttttt atttttatatt 60
tatttgagaa acccccaaatt ttgctctgat ggcctttctt ctccatttgt catctctggg 120
aaccttgagt ctagatttag ggctgaaagt ctctctgcta ggagctttcc aaaatgcatt 180
gctttgattc tgggaactga ataggaggag aacacctgga ctactctgag tcctgagttc 240
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ggggctggta gagctgggtc ttgggcaac a gagtctgctt gtctgnaagc tctcgactgg 360
gcgaactcca tctgtctcag caatgaagta gaccccaaag gcaaggac 408

<210> 415
<211> 457
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(457)
<223> 5' terminal sequence. cd3g antigen, gamma
polypeptide (tit3 complex) (CD3G) gene.

<400> 415
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gcagaagcca aaaatatcac atggttttaa gatgggaaga tgatcggctt cctaactgaa 120
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tcagacaagc agactctgtt gcccaatgac cagctctacc agccctcaa ggatccgagg 420
aagatgacca gtacagccac cttcaagggn aaccagt 457

<210> 416
<211> 822
<212> DNA/RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature

262/292

<222> (1)..(822)

<223> cd3g antigen, gamma polypeptide (tit3 complex) (CD3G) gene.

<400> 416

```

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```

<210> 417

<211> 439

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(439)

<223> 3' terminal sequence. v-rel avian
reticuloendotheliosis viral oncogene homolog b
(nuclear factor of kappa light polypeptide gene
enhancer in b-cells 3) (RELB) gene.

<400> 417

```

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gaaggtgccc ccacatgtgg aacagaggca gctgtaacaa gctagtgcac gggagccatg 180
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aggggatggg ggtgctagac atcctgggtt gggattgcac ggctcctcca cctccctccc 360
caccagtgcc cctcctctg gcacgcgggg gctacgtggc ttcaggcccg gggataggag 420
gccgccccca aagccgct

```

43 9

<210> 418

<211> 234

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(234)

<223> 5' terminal sequence. v-rel avian
reticuloendotheliosis viral oncogene homolog b

263/292

(nuclear factor of kappa light polypeptide gene
enhancer in b-cells 3) (RELB) gene.

<400> 418

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caaacggcgg aagaaaaagc cggccatcct ggaccacttc ctgcccaacc acggctcagg 120
cccgttcctc ccgcgcgcag ccctgctgcc agaccctgac ttctttcttg ggcaccgtgt 180
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```

<210> 419

<211> 2314

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2314)

<223> v-rel avian reticuloendotheliosis viral
oncogene homolog b (nuclear factor of kappa light
polypeptide gene enhancer in b-cells 3) (RELB)
gene.

<400> 419

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```

264/292

```
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cagacttgac ttgaagggtg ggggtaggtt ggttggtcag agtcttccca ataaagatga 2280
gtttttgagc ctcaaaaaaa aaaaaaggaa ttcc 2314
```

<210> 420

<211> 214

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(214)

<223> 3' terminal sequence. estrogen receptor 1
(ESR1) gene.

<400> 420

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gagcatcccg ctggattctt tttcaaagt caaaagaggt ttacaagtgt gtttcattaa 120
acaaagcaaa gctgcgacaa aaccgagtca catcagtaat agtatgcac ggcaaaaggg 180
catattaatc catcaaacac aatttgcat ttga 214
```

<210> 421

<211> 520

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(520)

<223> 5' terminal sequence. estrogen receptor 1
(ESR1) gene.

<400> 421

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gttttagagct gtcacctag aaaca acata ttgtcccatg agcaggtgcc tgagacacag 360
acccctttgc attcacagag aggtcattgg ttatagagac ttgaattaat aagtgcatt 420
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cagtgtagag ctottgtttt atgggaaaag gctcaaatgc 520
```

<210> 422

<211> 6450

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(6450)

<223> estrogen receptor 1 (ESR1) gene.

<400> 422

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266/292

```

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ctttccaatt gaattaaagt gtggcctcgt ttttagtcat ttaaaattgt tttctaagta 5880
attgctgcct ctatta tggc acttcaattt tgcactgtct tttgagattc aagaaaaatt 5940
tctattcatt tttttgcact caattgtgcc tgaactttta aaatatgtaa atgctgccat 6000
gttccaaacc catcgtcagt gtgtgtgttt agagctgtgc accctagaaa caacatactt 6060
gtccatgag cagggtgcctg agacacagac ccctttgcat tcacagagag gtcatt tgggt 6120
atagagactt gaattaataa gtgacattat gccagtttct gttctctcac aggtgataaa 6180
caatgctttt tgtgcactac atactcttca gtgtagagct cttgttttat gggaaaaggc 6240
tcaaagcca aattgtgttt gatggattaa tatgcccttt tgccgatgca tactattact 6300
gatgtgactc ggttttgtcg cagctttgct ttgtttaatg aaacacactt gtaaacctct 6360
tttgactttt gaaaaagaat ccagcgggat gctcgagcac ctgtaacaa ttttctcaac 6420
ctatttgatg ttcaataaaa gaattaaact
6450

```

<210> 423

<211> 580

<212> DNA

<213> Artificial Sequence

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<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(580)

<223> 3' terminal sequence. pre -b-cell leukemia transcription factor 1 (PBX1) gene.

<400> 423

```
ttaaagctac aaacactttt attattttct ttgtaatttt tttcctcttt aaattcctct 60
aattgttgaa aatatccttc agtgatatgc gagaggcgcg gcacccggga gtctaggaca 120
gaggcacagg ggcagggaag atgacgaaaa ccaggctgac agctggaggc aggaagggt 180
ggcttctacc cagaaaaaaaa aggggagaga gtataaagaa gtgtccagat tggctgaaat 240
agcatcccaa agaagagaag agaaggagac tcttattgtg ttgctgatt gcttcgacct 300
ccagtctgac cgcttcaggc ttgggagaga aaccctccct tcctgcccct gccccaactg 360
ggcgacaggg tcagccggga tgcgattgct gggagatcag ttggaggat cagagtgaac 420
actgccaggg ccttctgtag gggaggtcac tgatgaagg gtagtagcat cctgccaacc 480
tccattagca ctgatgcct gcg gactgta catctgactg gctgagagtc catcactgta 540
tcctcctgtc tggctgataa catggcgaag ggtatccacc 580
```

<210> 424

<211> 503

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(503)

<223> 5' terminal sequence. pre -b-cell leukemia transcription factor 1 (PBX1) gene.

<400> 424

```
gccaaagagg agttagccaa gaagtgtggc accacagtct ccaggtatc acaactggtt 60
tggaaataag cgaatccggt acaagaagaa cataggtaaa tttcaagagg aagccaatat 120
ttatgctgcc aaaacagctg tcaactgtac caatgtgtca gccatggaa gccaaagctaa 180
ctcgccctca actcccaact cggtgtgttg atacccttcg ccatgttatc agccagacag 240
gaggatacag tgatggactc gcagccagtc agatgtacag tccgcaggca tcagtgtctaa 300
tggagggttg caggatgcta ctacccttc atcagtgaac t cccctacag aaggccctgg 360
cagtgttcac tctgatacct ccaactgac tccagcaat cgcaccccg ctgaccctgt 420
gccccagttg ggcaggggca ggaggagggt tttctctccc aacgctgaag cggtcagact 480
ggagggtcaaa cgattagcca aac 503
```

<210> 425

<211> 1819

<212> DNA/RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(1819)

<223> pre-b-cell leukemia transcription factor 1 (PBX1) gene.

```

<400> 425
cttccctggt tatcctgaaa aggatttgaa g acaagcttg aaggataaaa agccttggtg 60
cttcccagga gccgagccga ggagcagaag aggaagagcc gggggctgcc gtagcctttg 120
gagatggacg agcagcccag gctgatgcat tcccatgctg gggcgggat gcccgacac 180
cccggcctgt cccagcactt gcaggatggg gccggaggga ccgaggggga gggcgggagg 240
aagcaggaca ttggagacat tttacagcaa attatgacca tcacagacca gagtttggat 300
gaggcgagg ccagaaaaa tgctttaaac tgccacagaa tgaagcctgc cttgtttaat 360
gtgttgtgtg aaatcaaaga aaaaacagtt ttgagtatcc gaggagccca ggaggaggaa 420
cccacagacc cccagctgat gcggctggac aacatgctgt tagcggaagg cg tggcgggg 480
cctgagaagg gcggagggtc ggcggcagcg gcggcagcgg cggcggttc tggaggggca 540
ggttcagaca actcagtggg gcattcagat tacagagcca aactctcaca gatcagacaa 600
atctaccata cggagctgga gaaatacgag caggcctgca acgagttcac caccacagt 660
atgaatctcc tgcgagagca aagccggacc aggccatct ccccaaagga gattgagcgg 720
atggtcagca tcatccaccg caagttcagc tccatccaga tgcagctcaa gcagagcacg 780
tgcgaggcgg tgatgatcct gcgttccga tttctggatg cgcggcgga gagacggaat 840
ttcaacaagc aagcgacaga aatcctgaat gaatatctt attcccatct cagcaaccct 900
taccocagtg aggaagccaa agaggagtta gccagaagt gtggcatcac agtctcccag 960
gtatcaaact ggtttgaaa taagcgaatc cgttacaaga agaacatagg taaatttcaa 1020
gaggaagcca atatttatgc tgccaaaaca gctgtcactg ctaccaatgt gtcagcccat 1080
ggaagccaag ctaactcgcc ctcaactccc aactcggctg gttcttcca g ttttttaac 1140
atgtcaaact ctggagattt gttcatgagc gtgcagtcac tcaatgggga ttcttaccaa 1200
ggggcccagg ttggagccaa cgtgcaatca caggtggata cccttcgcca tgttatcagc 1260
cagacaggag gatacagtga tggactcgca gccagtcaga tgtacagtcc gcagggcac 1320
agtgtaatg gaggttgga gga tgctact accccttcat cagtgcctc ccctacagaa 1380
ggccttgga gtgttctc tgatacctc aactgatctc ccagcaatcg catcccggct 1440
gacctctgc cccagttggg gcaggggcag gagggagggt ttctctccca agctgaagcg 1500
gtcagactgg aggtcgaagc aatcagcaaa cacaataaga gtctccttct cttctcttct 15 60
ttgggatgct atttcagcca atctggacac ttctttatac tctcttccct tttttttctg 1620
ggtagaagcc acccttccct gcctccagct gtcagcctg ttttgcctat cttccctgcc 1680
ctgtgcctc tgcctagac ttcccgggt ccccgccctc tctcatatca ctgaaggata 1740
ttttcaacaa tttagggaat ttaaagagga aaaaaat tac aaagaaaata ataaaagtgt 1800
ttgtacgttt tcaaaaaaa 1819

```

<210> 426

<211> 506

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(506)

<223> 3' terminal sequence. gli-kruppel family
member gli3 (greig cephalopolysyndactyly syndrome)
(GLI3) gene.

<400> 426

```

taaaaagagg gtggtttgag tgtaacaata ctgattcaaa actgaaatgg aagacagttt 60
ctccctagaa tactttaggg tttttcagag tctttttcca taaa aggaat ataattgaaa 120
cacatctcag ttaggtgaga tgagattgct aaaatacata cagaactaaa aaaacagcca 180
aaacaaagtc agtttaatct cttcaactcc taatgatttc cgttgggttc agtctttttt 240
tcttaaagcc tattgcataa ctgcaaggga attgctttct tccgctaggg aggtcagcaa 300
agaactcatg tccccgatag cc atgttggt ggtgctcatg gacagcgtg ggaatgggag 360
ggacgcccga gccgtggtta aagccggaag aacctatgga aaaggtctca atgatacttg 420
ggctcagggc cccgacatca ggctcgagt gtcccatct ctatgattgc atcgaagtca 480
atctggtacc ctcccaggcc aggctg 506

```


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<210> 427
 <211> 239
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(239)
 <223> 5' terminal sequence. gli-kruppel family
 member gli3 (greig cephalopolysyndactyly syndr ome)
 (GLI3) gene.

<400> 427
 ggcagaagga tcacttgagg ccaggcattc aagaccagcc tatgcaagat aatgagacct 60
 tgtctctatt taaaaaacia aaagcctggt gtggtggtgc atgcctgtag tctcagcctc 120
 ctgagtagct gggactataa gcacgcacca ctatgtctgg ctaatttttg tatttttagt 180
 agagacaggg ttccaccacg ttgcccaggg cagtcttgaa ctctgaccc caagtgate 239

<210> 428
 <211> 5054
 <212> DNA/RNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(5054)
 <223> gli-kruppel family member gli3 (greig
 cephalopolysyndactyly syndrome) (GLI3) gene.

<400> 428
 cgatactacg tgggcatttt tggcgaaga gagctgaagt aatgagaaga catcatggag 60
 gcccagtcac acagctccac gaccactgaa aagaaaaaag ttgagaattc catagtgaag 120
 tgcctccactc gaacagatgt gagcgagaaa gccgttgctt ccagcaccac ttctaattgag 180
 gatgaaagtc ctggacagac ttatcacaga gagagaagaa acgcaatcac tatgcagcca 240
 cagaatgtcc aggggctcag caaagtcagt gaggaacctt caacatcgag tgacgagagg 300
 gcctcattga tcaagaaaga gatccatggg tccctgccac acgtggcgga gccctctgtg 360
 ccgtaccgag ggacggtg tt tgccatggac cccaggaatg gttacatgga gcccactac 420
 caccctcttc atcttttccc tgccttccat cctcctgtac caattgatgc cagacatcat 480
 gaggggcgtt accattacga tccatctccg attcctc
 cat tgcataatgac ttccgcctta 540
 tctagtagcc ctacgtatcc ggacctgcc ttcattagga tctccccaca ccggaacc cc 600
 gctgctgctt ccgagtctcc cttcagccct ccacatccct acattaatcc ctacatggac 660
 tatatccgct ccttgcaacag cagcccatcg ctctccatga tctcagcaac ccgtgggctg 720
 agccctacag atgcgcccc tgcaggagtc agcccagcag aatactatca tcagatggcc 780
 ctgctaactg gccagcgag cccctatgca gacatt attc cctcagctgc caccgcccgc 840
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 aggtgttcag ccaggccgag ccgaaaacgt acactgtcca tatcaccact ctccgatcat 960
 agctttgacc ttcagaccat gataaggagc tctcccaact ccttggtcac gattctcaat 1020
 aattcccgta gcagctcttc agcaagtggc tccatgggtc acttatctgc aagtgcaatc 1080
 agccctgctt tgagcttcac ctactcttcc ggcgccgtct ctctccacat gcatcagcag 1140
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 gagcagaaac ccttcaaagc ccagtatatg ttggtagtgc atatgagaag acacacgggc 1680
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 aatgagaaac catatgtgtg caaaatccca ggctgcaact agcgttacac agaccaagc 1920
 tccctccgga aacatgtgaa gacagtgcac ggcccagagg ctcatgtcac caagaagcag 1980
 cgaggggaca tccatccctg gccgccaccc ccgagagatt ccggcagcca ttcacagtcc 2040
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 cagattgact tcatgacct catagacgat ggggaccact ccagcctgat gtcgggggccc 4620
 ctgagcccaa gtatcattca gaacctttcc catagctcct cccgcctcac cagcctcgg 4680
 gcgtccctcc cattccagc gctgtccatg ag caccacca acatggctat cggggacatg 4740
 agttctttgc tgacctccct agcggaagaa agcaaatcc ttgcagttat gcaataggct 4800
 ttaggaaaaa aagactgcaa ccaacggaaa tcaataggag ttgaagagat taaactgact 4860
 ttgttttggc tgttttttta gttctgtatg tattttagca atctcatctc acctactga 4920

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gatgtgtttc aattatattc cttttatgga aaaggactct gaaaaaccct aaagtattct 4980
 agggagaaac tgtcttccat ttcagttttg aatcagtatt gttacactca aaccaccctc 5040
 tttttaaaaa aaaa 5054

<210> 429
 <211> 271
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(271)
 <223> 3' terminal sequence. interleukin enhancer
 binding factor 1 (ILF1) gene.

<400> 429
 aggagacttg ttttactcag agtggaaaat tttgccaggg acaaagtc aa cacaaagaaa 60
 caaacaacaa aaaatagcca gaaagagaac agttaagtgc agctcgggtga gtcccggcag 120
 ttccttcccg gcactggctc gtccctgggt tctcaagggt ccatgcggcc acagcgccg 180
 tccacctgtc cagcgcagcc acatgctgaa atggagggtg ataaaattca tcaggcagct 240
 gctgtaacac ggaaatgtgc agatgcc aga g 271

<210> 430
 <211> 193
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(193)
 <223> 5' terminal sequence. Similar to Retinoblastoma -like protein 2,
 sw|Q08999|RBL2_HUMAN.

<400> 430
 TGTCTACATT TNCCACATA AAAATGAAAC AATNNTTCT CTCGNGAAA AGATTTTATA 60
 TTA CTNAGC NAAAGTCCTT CAAAGNGACT GAGCGNANNT AATAGTNTGA TACGCATAGG 120
 NGNAACTCCT ACTNNAANGN GCGGNNTTCT TNTNGNGAT GGCNGTGAAT NCACTGCAAA 180
 NAGCATTTGC CCN 193

<210> 431
 <211> 184
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(184)
 <223> 3' terminal sequence. Homo sapiens CD2 antigen (p50).

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<400> 431
AGTCAAAGCT TTTATTTAC TTTNACTCAC AGGATGGGGG GCAAGTNTCC ACCAGGGCAC 60
ATAAGAAACT CCAGAGTCTC TTAAGCAGAT AGGCTGCTTG TAGNGAGACC CAGGNACACC 120
AATCACTTGA TTTATCGCAT CTACACATGA CCNGAGAGGG GACAAGAAAT CTCTAAGTTT 180
TCTG 184

<210> 432
<211> 242
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(242)
<223> 5' terminal sequence. Similar to NM_022641 Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1)

<400> 432
ATTGGGCACG ATGACTGGCC AGGGTATAAA AAGGGCCAC AAGAGACCGG CTCTAGGATC 60
CCAAGGCCCA ACTCCCGAA CCACTCAGGG TCCTGTGGCA GCTCACCTAG TGGCAATGGC 120
TCCAGGCTCC CGGAACGTCC CTGATCCTGG NTTTTGNCCT NCTCTNACTG CCCTGGNTTN 180
AANAAGCTNG TGCCNTCCAA ANCGTTCCTG TATCCAGGGT TTTTGACCAC GCTATGCTNC 240
AA 242

<210> 433
<211> 329
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(329)
<223> 5' terminal sequence. Homo sapiens plasminogen activator (PLAT)

<400> 433
AACAGTCACC GACAACATGC TGTGTGCTGG AGACACTCGG AGCGGCGGGC CCCAGGNAAA 60
CTTGACGAC GCCTGCCAGG GNATTCGGG AGGCCCCCTG GTGTGTCTGA ACGATGGCCG 120
CATGACTTTG GTGGGCATCA TCAGCTGGGG CCTGGGCTNT GGACAGAAGG ATGTCCCGGG 180
TGTGTACACC AAGGTTACCA ACTACCTAGA CTGGATTCGT GACAACATGC GACCGTGACC 240
AGGAACACCC GACTCCTCAA AAGNAAATGA GATCCCGNCT CTTCTTCTTC AG AAGACACT 300
GCAAAGGCGC AGTGCTTCTC TACAGACTT 329

<210> 434
<211> 247
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature

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<222> (1)..(247)

<223> 3' terminal sequence. Homo sapiens plasminogen activator (PLAT)

<400> 434

AAATATATCT GAANNNTTAA AGTACAGTNT AAAACAGGGT TNTGGCAACA GAAAGTAAAA 60
ACTAACATGG NTTGCTATAA ATATGCTGAA GCCTAGTTGT TCAAATGATA CAATTCTCTC 120
ATGCTACTCT AAAGTTTNTA AAGNAAAAGG GTTTACACTT TACACACT GT ACACAAAGGG 180
GNTACCTTCT GNGNGCCNGG GAGTGGGGAA AGGGGANGSN GACTTGACGT CAAGGGTGCT 240
TTTGAGG 247

<210> 435

<211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(63)

<223> 5' terminal sequence. Homo sapiens aminoacylase 1 (ACY1).

<400> 435

CCGCCAGNAC CTGCGTATCC GCACTGTTCA GCCAAGCCT GACTATGGAG CTGCTGTGGC 60
TTT 63

<210> 436

<211> 190

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(190)

<223> 3' terminal sequence. Homo sapiens aminoacylase 1 (ACY1).

<400> 436

GAAGAGGAGG NCCTTGGCAC TGGTNGGGAN GGAAGCCCCA GGGGAAAAGG TTNAGGAGTC 60
CCAGGGCTCA GCTGTCACTG GGCAGGGCAG GCACACTGGC AGGGCCAGGC AGCAGGCNNG 120
TATATATGNC CACCCACGG AGGAACACAG CCTNATGCAG NCGTTNATCG TGGTNGTGCA 180
GNAGCACAGG 190

<210> 437

<211> 176

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(176)

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<223> 3' terminal sequence. Homo sapiens E74 -like factor 1 (ets domain transcription factor) (ELF1)

<400> 437

CCCTTAGGCT GTTCTGGTGA TGAGGCTCCC GGTGAGTCTG CATATTTTNN TTGCACCTGC 60
TGTGTTTCCA TCACTTCAGG AATCCCATCT AATGTGACGG ACACATGGGT GACTGGGGCA 120
ACAACCATGT CATCTTCAGG NGAACATAAT ATATTATTAT TTATCCGGTT TTCATC 176

<210> 438

<211> 465

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(465)

<223> 5' terminal sequence. Homo sapiens selectin P,
granule membrane protein CD62, SELP

<400> 438

NGGAACATAT GGAGTTTTTA CA AACGCTGC ATTTGACCCG AGTCCTTAAG AGACCTGTCC 60
TTTTCTGGT CTCCTCATTC AGCCTCCATA TGATCCTGTT GTGAACATCA AGTTTCCTGC 120
TACACTGGAC TTAACATAAA TGCATTTGCT GCAGGTTTCC ATAAACACCC ATGANTCAAA 180
GACATGGAAT TACCTTAGAT TAGCTCTGGA CCAGCCTGTT GGACCCGCTC TGGACCAACC 240
CTGTTTCCTG AGTTTGGGAT TGTGGTACAA TCTCAAATTC TCAACCTACC ACCCCTTCCT 300
GTCCACCTC TTCTCTTCCT GTAACACAAG CCACAGAAGC CAGGAGCAA TGTTCCTGCA 360
GTAGTCTCTG TGCTTTGACT CACCTGTTAC TTGAAATACC AGTGAACCAA AGAGACTGGA 420
GCATCTGGAC TTNACAAGAA GACCAGACTT GTGGAGANAT TAA AA 465

<210> 439

<211> 396

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(396)

<223> 3' terminal sequence. Homo sapiens selectin P,
granule membrane protein CD62, SELP

<400> 439

GAAAATCCTT TNAATTACGN ATTTNAANAT TGGNCTTTGG GNCATTTGAG GGACAGTGAC 60
TGGGGGCAGG AAGGCCACAG GNGACATGTG GCCTTCTAGC TTNATNCTTG GCCTTCTGCA 120
GCCTCTGGGG CCATCCAGAG GACTCTCTGG AAGCCTCAGA GCAGAGGNCC AAGAGGCCCTC 180
ATCAACAGCA ACCANTGGAG ACTCCAGAAG ATGCTACAGG ANTTNCTCCC AATTAGAGAT 240
GCCACCTGTT TNCCAACAAA GTGGAGAAAA CCTTCCTNNA ATCAAAAANT AAAGAGGTNT 300
TTCNATTTCT CCACAGTTTG GCCTCCTTGT GAGTCAGATG NTCCAGTCTC TTTGGNNCAC 360
TGGTATTTCA AGTAACAGGT GAGNNAAGG ANAGAG 39 6

<210> 440

<211> 337

<212> DNA

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(337)

<223> 5' terminal sequence. Human interleukin 3 receptor (hIL -3Ra).

<400> 440

```
AAAGGCTCAG CAGTTGAC CT GGGACCTTAA CAGAAATGTG ACCGATATCG AGTGTGTTAA 60
AGACGCCGAC TATTCTATGC CGGCAGTGAA CAATAGCTAT TGCCAGTTTG GAGCAATTTT 120
CTTATGTGAA GTGACCAACT ACACCGTCCG AGTGGCCAAC CCCCATTCT CCACGTGGAT 180
CCTCTTCCCT GAGAACAGTG GGAAGCCTTT GGCAGGTGCG GAGAATCTGA CCTGCTNGAT 240
TCATGACGTT GATTTCTTGA GCTGCAGCTG GCGGGTAGGG CNGGGGGCCC CGCGGGACGT 300
CCAGTACGAC CTGTAATTGA ACGTNNCCAA CAGNGT 337
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<210> 441

<211> 104

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:prim er

<220>

<221> misc_feature

<222> (1)..(104)

<223> 3' terminal sequence. Human interleukin 3 receptor (hIL -3Ra).

<400> 441

```
TTCCCGCCTC CCAGACCACC AGNTTGNCGT TTTGGNAGNT NTNACCGANG GGGTNTTNCA 60
TGTNAGGGNT NCGGNAAN AGTTTNTGAA NCACCAGAAA CCTT 104
```

<210> 442

<211> 223

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(223)

<223> 5' terminal sequence. Human tumor protein p53 (Li -Fraumeni syndrome) (TP53)

<400> 442

```
AAGCAATGGA TGATTTGATG CTGTCCCCGG ACGATATTGA ACAATGGTTC ACTGAAGACC 60
CAGGTCCAGA TGAAGCTCCC AGAATGCCAG AGNTGNCTCC CCGCGTGGCC CCTGCACCAG 120
CAGCTCCTAC ACCGGCNGCC CCTGCACCAG CCCCTCCTG GCCCTGTCA TCTTCTGTCC 180
CTTCCAGAN AACCTACCAG GGCAGCTACG GTTCCGTCT G GG 223
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<210> 443

276/292

<211> 223
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(223)
<223> 5' terminal sequence. Tumor protein p53
(Li-Fraumeni syndrome) (TP53) gene.

<400> 443
aagcaatgga tgatttgatg ctgtccccgg acgatattga acaatgggtc actgaagacc 60
cagggtccaga tgaagctccc agaatgccag agntgnctcc ccgcgtggcc cctgcaccag 120
cagctcctac accggcngcc cctgcaccag cccctcctg gccctgtca tcttctgtcc 180
cttcccagan aacctaccag ggcagctacg gtttccgtct ggg

<210> 444
<211> 343
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(343)
<223> 3' terminal sequence.

<220>
<221> misc_feature
<222> (1)..(343)
<223> v-erb-b2 avian erythroblastic leukemia viral
oncogene homolog 2 (neuro/glioblastoma derived
oncogene homolog) (ERBB2) gene.

<400> 444
caaggggctg caaactnncc cacacatgac ccagccctc tacagcggta cagtgaggac 60
cccacagtac cctgcccctc tgagactgat ggctacgttg cccccctgac ctgcagcccc 120
cagcctgaat atgtgaacca gccagatgtt cggccccagc ccccttcgcc ccgagaggcc 180
ctctgcctgc tgcccgaact gctggtgccca ctctggaaag gcccaagact ctctccccag 240
ggaagaatgg ggtcgtcaaa gagtttttgc cttt ggggggt gccgtgggag aaccccagat 300
attgacaccc caggggaggg agcttgccct tcagcccccac ctt 343

<210> 445
<211> 309
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(309)
<223> 3' terminal sequence.

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<220>
<221> misc_feature
<222> (1)..(309)
<223> zinc finger protein 144 (Mel -18) (ZNF144) gene.

<400> 445
ccgcccccca cccccaaggt gganagagct ggggaaagta gaagaggtgg aaaaaagggc 60
ccagaaaaag tggaaggagt ggagaggctt ag ctggaaga agggagaggg tccctggcct 120
caagttaagg ggggcacggg agcgccnttg acagtcatct tgcgccccct gctggtingaa 180
ganggtttct tcaggcagtt caagctaccc ccgttingcan ctttngnngc ccacttgct 240
ntcgaagggg ganttgggna ngtagggting gtngcttgan gcccatngga actnggaaaa 300
ccatnggat 309

<210> 446
<211> 268
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(268)
<223> 3' terminal sequence.

<220>
<221> misc_feature
<222> (1)..(268)
<223> MAP/microtubule affinity -regulating kinase 3
(MARK3) gene.

<400> 446
attatcttgt gaatctactt agaaaaacac acacaagcaa tggtcacaac tataaattta 60
aaccttttgc actaaaaaaa cacaaaacaa caaacacaan accacaggca tgaactgtaa 120
acctgtatta actntgaact ggtcttaagg ttaattctta gcngtcattc agtattttcc 180
tccttggtcaa ctgtaatgtt ttngcaccgg ntgatctccc gcnggggggta ctagtaatga 240
ctggctgccc gtgtagggag atgcttcc 268

<210> 447
<211> 169
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(169)
<223> 5' terminal sequence.

<220>
<221> misc_feature
<222> (1)..(169)
<223> MAP/microtubule affinity -regulating kinase 3
(MARK3) gene.

<400> 447

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gaacactgat ggaaatgtat agaataatat ttaggcaata acgtctgcat cttctaaatc 60
 atgaaattaa agtctgagga cgagagcacg nctggngcg aaantntgcc tttttntac 120
 ggatgcacta cantaaagan gtgcanccta tncgnccct gccctactt 169

<210> 448

<211> 393

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(393)

<223> 3' terminal sequence.

<220>

<221> misc_feature

<222> (1)..(393)

<223> EST N68536 MAX-interacting protein 1 (MXI1) gene.

<400> 448

aagtaattaa gggtagttaa attatttaaa gtatacaaag tccaaacagc caggggtaag 60
 gtctccaaga ggccttccca gggtaaagga gagggccaca agggaacctt ggagtttgaa 120
 agacaaaggg aacacatgac atcaaagtgc aggctagaaa ttctacttag aagaaaataa 1 80
 cattactgaa aatattcgta agagtaacaa tagcacatgc acagtgggac tgtgaggaag 240
 agagagactg cctgtaggaa aatggaagca aatctttaca ttaaaatgag acaagtgctg 300
 aacttactat gttaactatg atagtgtgtc tacnatagat atcnogatgg ttaaaagctg 360
 gtaaaaggta atggttctca aaacnnaaa a tag 393

<210> 449

<211> 217

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(217)

<223> 5' terminal sequence.

<220>

<221> misc_feature

<222> (1)..(217)

<223> EST R81126 lymphotoxin beta receptor (LTBR) gene.

<400> 449

ttacntgggt atctectact gtagtatgag gaagaatggc tgtaaatgta ttttttgaat 60
 tctggnctca cctttgtctc agctaaatgt agccgcaccc gggacacagt ttgtgccaca 120
 tgtgccgaga attcctacaa cgagc actgg aactacatga ccatctgccca nctgtgccgc 180
 ccctgtgacc cagtgatggg cctcgnctga gantgcc 217

<210> 450

<211> 157

<212> ADN

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(157)

<223> 5' terminal sequence

<220>

<221> misc_feature

<222> (1)..(157)

<223> (POU2F2) gene.

<400> 450

nattcggcaa cgnggaaagg aaagaaacta accaacaaaa gagaaaacca aaaataatca 60
caacagaaac cagctgcccc aaaggaggcc agtngtnggg acgcagaggg tcc tcagagc 120
aggagtnaca agggaggaaa gaccaaaaaa acaacca 157

<210> 451

<211> 282

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(282)

<223> 5' terminal sequence.

<220>

<221> misc_feature

<222> (1)..(282)

<223> caspase 4, apoptosis -related cysteine protease
(CASP4) (ex CASP1)

<400> 451

gagaatctga cagccaggga tatggagtca gcgctgaggg catttgctac cagaccagag 60
cacaagtctc ctgacagcac attcttggtg ctcatgtctc atggcatcct ggagggaatc 120
tgcggaactg tgcatgatga gaaaaaacca gatgtgctgc tttatgacac catcttccag 180
atattcaaca accgcaactt gcctcagtct ngaaggacaa acccaaggtc atcattgtnc 240
agggccttgca gaggttgcaa accttggggg aantttgggg tc 282

<210> 452

<211> 424

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(424)

<223> 5' terminal sequence

<220>
 <221> misc_feature
 <222> (1)..(424)
 <223> syndecan 1 (SDC1) (ex HRB)

<400> 452
 ttgcttttng caaaactcta cttaatccaa tnggtttttc cctgtacagt agattttcca 60
 aatgtaataa actttaatat aaagtagtcc tgtgaatgcc actgccttcg cttcttgect 120
 ctgtgctgtg tgtgacgtga cgggactttt ctgcaaacac caacatgttg ggaaacttgg 180
 ctogaatctc tgtgccttcg tctttcccat ggggagggga ttctgggtcc agggccctc 240
 tgtgtatttg cttttttgtt ttggctgaaa ttctcctgga ngtcggtagg ttcagccaag 300
 gttttataag ggctgatgtc aatttntctgt gtttgccaan ttccaagccc catcttncta 360
 aatgggcaaa aggaaggtgg gatgggcccc agcnacagct ttgnaccc ga ggctnttgg 420
 gtca 424

<210> 453
 <211> 435
 <212> ADN
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(435)
 <223> 5' terminal sequence

<220>
 <221> misc_feature
 <222> (1)..(435)
 <223> Itegrin, beta 2 (antigen CD18 (p95), lymphocyte
 function-associated antigen 1; macrophage antigen
 1 (mac-1) beta subunit) (ITGB2)

<400> 453
 aggagtgcgc cggctgcccc tcaccct gtg gcaagtacat ctccctgcgc gagtgccctga 60
 agttcgaaaa gggccctnt ggaagaactg cagcgcggcg tgtccgggcc tgcagctgtc 120
 gaacaacccc gtgaagggca ggacctgcaa ggagaggac tcagagggt gctgggtggc 180
 ctacacgtg gagcagcagg acgggatgga ccgctacctc atctatgttg atgagagccg 240
 agagtgtgtg gcaggcecca acatcgccgc catcgtcggg ggcaccgtgg gcaggcatcg 300
 tgctgatcgg catctcctc ctggtcatct gggaaggctc tgatccacct gagcgacctc 360
 cgggagttac aggcgttttg agnaggagaa gctcaagtcc cagtnggaac aatgattatt 420
 ccctttttca agagc 435

<210> 454
 <211> 544
 <212> ADN
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(544)

<223> 5' terminal sequence

<220>

<221> misc_feature

<222> (1)..(544)

<223> Protein phosphatase 1, catalytic subunit, alpha
isoform (PPP1CA) (ex MGST1)

<400> 454

```
gtgtgaccag acatgcaacc gncatctatg gtttctacgn atgnagtgn aagcagnacg 60
nctnacaaca tcaaactgtg gnaaaacctt cactgnactg ncttcaactg ncctgnccca 1 20
tcgcggncca tagtggacgt aaaagatctt ctgnctgncc acggaggcct gttccccgga 180
cctgncagtt ctatggnagc agattcggcg ggatcatgcg gcccacagat gtgtcctgta 240
ccaggggcct gctgtgtgac ctgctgtggt ctgacctga caaggacgtg cagggtctggg 300
gcgagaacga cgtgtggcgt ctcttttac c ttggagccg aggtggtggc caagttcctc 360
cacaagcacg acttggacct catctgccga gcacaccagg ttgtagaaga cggctacgag 420
ttctttgcc aagcgcantt ggtgacactt ttctcagctt ccaactactg ttgcgagttt 480
nacaatgctt gcgccatgat gagtgtgacg agacctatg tgcttttcag atcttaagcc 54 0
gccn 544
```

<210> 455

<211> 344

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(344)

<223> 3' terminal sequence

<220>

<221> misc_feature

<222> (1)..(344)

<223> protein phosphatase 2 (formerly 2A), catalytic
subunit, alpha isoform (PPP2CA)

<400> 455

```
actgcggtga gagccagcgg ccagcgccac cncaacagcc gccagaagna cagcaggaac 60
cggcggcggc gngtgcngt aggcccggtg n gcggtgncgg cgcgggagga gccggannca 120
gccggcnggg gcggnggca ncanggacga gaaggngnnc accaaggagc tggaccagng 180
gaacgagcag cngaacgagn gcaagcagcn gnccgagncc caggncaga gccncagcga 240
gaangcnaaa gaaanccnga caaaagaanc caacngcaa gaggnncgan gnccaggnac 30 0
ngnccngnga gangngcang ggcaannnca ngaacccatg gaac 344
```

<210> 456

<211> 514

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(514)

<223> 5' terminal sequence

<220>
 <221> misc_feature
 <222> (1)..(514)
 <223> S100 calcium-binding protein A11 (calgizzarin)
 (S100A11)

<400> 456
 cagcctcccg cgctcgctc agctccaaca tggcaaaaat ctccagccct acagagactg 60
 agcgggtgcat cgagtccttg attgctgtct tccagaagta tgctggaa ag gatgggttata 120
 actacactct ctccaagaca gagttcctaa gcttcatgaa tacagaacta gctgccttca 180
 caaagaacca gaaggaccct ggtgtccttg accgcatgat gaagaaactg gacaccaaca 240
 gtgatggta gctagatttc tcagaatttc ttaatctgat tggtaggcta gctatggctt 300
 gccatggact ccttcctcaa ggctgtccct tcccagaagc gggacctgga gggacccctt 360
 gggccctggg cctttcaaac ccacccctn ttcctttcca gcctttctgt tcatcatntt 420
 ccacagccca cccttncctg gaggcacatt aaccacctna tggtagggtn ccaactgggc 480
 attagtatt aaaggnaatg tnaatttttt ttaa 514

<210> 457
 <211> 359
 <212> ADN
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(359)
 <223> 5' terminal sequence

<220>
 <221> misc_feature
 <222> (1)..(359)
 <223> Granzyme A (granzyme 1, cytotoxic
 T-lymphocyte-associated serine esterase 3) (GZMA)

<400> 457
 gctggacgtc atcaacaagc attcattcaa caacttcgc ctgcgagtgg ggttgaacca 60
 tggaccgta gtagctggag ttattggggc ccagaagcog caatatgaca tttggggcaa 120
 cacagtgaac gtggccagcc gcatggagag tacaggagtc cttggcaaaa tccaagtgaac 180
 tgaggagaca gcatgggccc tacagtcctt gggctacacc tgctacagcc nggggtgtca 240
 tcanggtgaa aggcaaagg cagctctgca cctacttctt gaacacagac ttgacacgaa 300
 ctggacctcc ttcagctacc ctaggctgag attgcactcg cct tncctaag aacctnaat 359

<210> 458
 <211> 1251
 <212> ADN
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:primer

<220>
 <221> misc_feature
 <222> (1)..(1251)
 <223> 3' terminal sequence

<220>

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<221> misc_feature
<222> (1)..(1251)
<223> endothelin 1 (EDN1)

<400> 458
ggagctgttt acccccactc taataggggt tcaatataaa aagccggcag agagctgtcc 60
aagtcagacg cgctctgca tctgcccag gcgaacgggt cctgcgcctc ctgcagtccc 120
agctctccac caccgcgcg tgcgcctgca gacgtccgc tcgctgcctt ctctcctggc 180
aggcgtgcc ttttctcccc gttaaagggc acttgggctg aaggatcgct ttgagatctg 240
aggaaccgc agcgctttga gggacctgaa gctgtttttc ttcgttttcc tttgggttca 300
gtttgaacgg gaggtttttg atcccttttt ttcagaatgg attatttgct catgattttc 360
tctctgctgt ttgtggcttg ccaaggagct cc agaaacag cagtcttagg cgctgagctc 420
agcgcgggtg gtgagaacgg cggggagaaa cccactccca gtccaccctg gcggtccgc 480
cgggtccaagc gctgctcctg ctgcctcctg atggataaag agtgtgtcta cttctgccac 540
ctggacatca tttgggtcaa cactcccag cactgtgttc cgtatggact tggaagccct 600
agggtccaaga gagccttgga gaatttactt cccacaaagg caacagaccg tgagaataga 660
tgccaatgtg ctagccaaaa agacaagaag tgctggaatt tttgccaagc aggaaaagaa 720
ctcagggtcg aagacattat ggagaaagac tggaataatc ataagaaagg aaaagactgt 780
tccaagcttg ggaaaaagtg tatttatcag cag ttagtga gaggaagaaa aatcagaaga 840
agttcagagg aacacctaa acaaaccagg tcggagacca tgagaaacag cgtcaaataca 900
tcttttcatg atcccaagct gaaaggcaag ccctccagag agcgttatgt gaccacacaac 960
cgagcacatt ggtgacagac ttccgggctt gtctgaagcc atagcctcca cggagagccc 1020
tgtggccgac tctgcaactc ccaccctggc tgggatcaga gcaggagcat cctctgctgg 1080
ttcctgactg gcaaaggacc agcgtcctcg ttcaaaacat tccaagaaag gtaaggaggt 1140
tcccccaacc atcttcaactg gcttccatca gtggttaactg ctttgggtctc ttctttcatc 1200
tggggatgac aatggacctc tcagcagaaa caca cagtca cattcgaatt c 1251

<210> 459
<211> 2145
<212> ADN
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:primer

<220>
<221> misc_feature
<222> (1)..(2145)
<223> 3' terminal séquence

<220>
<221> misc_feature
<222> (1)..(2145)
<223> Protein tyrosine phosphatase, non -receptor type 6
(PTPN6)

<400> 459
cggcagaact gggaccaccg ggggtggtga ggcggcccg cactgggagc tgcattctgag 60
gcttagtccc tgagctctct gcctgccag actagctgca cctcctcatt ccttgccccc 120
ccttctctc cggaagcccc caggatgggtg agtggtgttc accgagacct cagtgggctg 180
gatgcagaga ccctgtctca gggccgaggt gtccacggta gcttcttggc tcggcccagt 240
cgcaagaacc aggggtgactt ctgcctctcc gtccagggtg gggatcaggt gacccatatt 300
cggatccaga actcagggga tttctatgac ctgtatggag gggagaagt t tgcgactctg 360
acagagctgg tggagtacta cactcagcag cagggtgtgg tgcaggaccg cgacggcacc 420
atcatccacc tcaagtaccc gctgaactgc tccgatecca ctagtgagag gtggtaccat 480
ggccacatgt ctggcgggca ggcagagacg ctgctgcagg ccaaggcgga gccctggacg 540
tttcttgtgc gtgagaacct cagccagcct ggagacttcg tgctttctgt gctcagtgac 600
cagcccaagg ctggcccagg ctccccgctc aggttcaccc acatcaaggt catgtgcgag 660
ggtggacgct acacagtggg ttggtttggag accttcgaca gcctcacgga cctgggtggag 720

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catttcaaga agacgggggat tgaggaggcc tcaggcgccct ttgtctacct gcggcagccg 780
tactatgccca cgagggtgaa tgcggtgac attgagaacc gagtgttgga actgaacaag 840
aagcaggagt ccgaggatac agccaaggct ggcttctggg aggagtttga gagtttgcag 900
aagcaggagg tgaagaactt gcaccagcgt ctggaagggc aacggccaga gaacaagggc 960
aagaaccgct acaagaa cat tctcccttt gaccacagcc gagtgatcct gcagggacgg 1020
gacagtaaca tccccgggtc cgactacatc aatgccaaact acatcaagaa ccagctgcta 1080
ggccctgatg agaacgctaa gacctacatc gccagccagg gctgtctgga ggccacggtc 1140
aatgacttct ggcagatggc gtggcaggag aacagccgtg tcatcgtcat gaccacccga 1200
gagggtggaga aaggccggaa caaatgcgtc ccatactggc ccgagggtggg catgcagcgt 1260
gcttatgggc cctactctgt gaccaactgc ggggagcatg acacaaccga atacaaactc 1320
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<210> 460

<211> 2149

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(2149)

<223> 3' terminal sequence

<220>

<221> misc_feature

<222> (1)..(2149)

<223> Transcription factor AP -4 (activating enhancer binding protein 4) (TFAP4)

<400> 460

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caccagggca gccgagagac ctccctcccg cccctcccat gccgcctcc ctccctcgc 120
cgccgcgcc gccgccagca tctgggaccg gccgattctg cactccgctc cgccgctgcc 180
ctttgattcg gatctccatc ttgcattctc cggctgatcg cgggacctgg ct cgtgcaga 240
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gaaggcatag gctccccgga catctgggag gacgagaagg cggaggacct gcggcgggag 720
atgattgagc tgcggcagca gctggacaag gagcgtcgg tgcgcatgat gctggaggag 780
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<210> 461

<211> 6478

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(6478)

<223> 3' terminal sequence

<220>

<221> misc_feature

<222> (1)..(6478)

<223> Cyclin D2 (CCND2)

<400> 461

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gctctccctt ccccttccaa aaaacaaaaa cagaaaaacc cttttccagg ccggggaaag 240
caggagggag agggcccgcc gggctggcca tggagctgct gtgccacgag gtggaccg 300
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tccaacctta catgcgcaga atggtggcca cctggatgct ggaggtctgt gaggaacaga 480
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<210> 462

<211> 3490

<212> ADN

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<220>

<221> misc_feature

<222> (1)..(3490)

<223> 3' terminal sequence

<220>

<221> misc_feature

<222> (1)..(3490)

<223> Junction plakoglobin (JUP)

<400> 462

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<210> 463

<211> 1355

<212> ADN

<213> Artificial Sequence

<220>

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 in (NME1)

<400> 464

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ccagatgggg tccagcgggg tcttggtgga gagattatca agcgttttga gcagaaagga 180
ttccgccttg ttggtctgaa attcatgcaa gcttccgaag atcttctcaa ggaacactac 240
gttgacctga aggaccgtcc attctttgac ggccgtggtg aatacatgca ctcagggccg 300

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290/292

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<211> 755

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 agaagagcag cctgtctg agaaaggcagt gaccaaggag gaatttcagg gtgaatggac 780
 tgcctccgct cctgagttca ctgctactca gcctgaggtt gcagactggt ctgaaggtgt 840
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<223> 3' terminal sequence

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lactogen) = LACTOGEN Precursor

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tcaggtcc cggaacgtcc ctgacctgg ntttgnct nctctnactg cctggnttn 180
aanaagctng tgccntccaa ancgttcgt tatccagggt tttgaccac gctatgctnc 240

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[FR/FR]; 971, chemin du Tardinaou, F-13190 Allauch
(FR).

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(74) Agents: **BREESE, Pierre** et al.; Breese-Majerowicz, 3,
avenue de l'Opéra, F-75001 Paris (FR).

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GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, PH, PL, PT, RO, RU, SD, SE, SG, SI,
SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU,
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GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent
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(71) Applicants (*for all designated States except US*): **IPSO-
GEN** [FR/FR]; 232 Boulevard Sainte-Marguerite, F-13273
Marseille Cedex 09 (FR). **INSERM** [FR/FR]; 101, rue
de Tolbiac, F-75654 Paris Cédex 13 (FR). **INSTITUT
PAOLI-CALMETTES - IPC** [FR/FR]; 232 Boulevard
Sainte Marguerite, F-13273 Marseille Cedex 9 (FR).

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(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **BERTUCCI,
François** [FR/FR]; Le Sully, Parc de la Cadenelle, 122,
rue du Commandant Rolland, F-13008 Marseille (FR).
HOULGATTE, Rémi [FR/FR]; 31, Boulevard Lom-
bard, F-13015 Marseille (FR). **BIRNBAUM, Daniel**
[FR/FR]; 9, rue Baldacchini, La Croix-du-Sud, F-13009
Marseille (FR). **NGUYEN, Catherine** [FR/FR]; 8 Boule-
vard de la Kabylie, F-13016 Marseille (FR). **VIENS,
Patrice** [FR/FR]; La Palmeraie Borely, Entrée A, 70,
avenue d'Haïfa, F-13008 Marseille (FR). **FERT, Vincent**

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*For two-letter codes and other abbreviations, refer to the "Guid-
ance Notes on Codes and Abbreviations" appearing at the begin-
ning of each regular issue of the PCT Gazette.*

(54) Title: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS USING ARRAYS OF CANDIDATE
GENES

(57) Abstract: The invention relates to a polynucleotide library useful in the molecular characterization of a carcinoma, the library including a pool of polynucleotide sequences of subsequences thereof wherein the sequences of subsequences are overpressed in tumor cells, further wherein the sequences of subsequences correspond substantially to any of the polynucleotide sequences set forth in any of SEQ ID NOS: 1-468 or the complement thereof. The invention relates also to polynucleotide arrays useful to differentiate tumor cells from normal cells comprising combinations of selected immobilized polynucleotide sequences sets.

WO 02/046467 A3

INTERNATIONAL SEARCH REPORT

Int Application No

PCT/IB 01/02811

A. CLASSIFICATION OF SUBJECT MATTER
IPC 7 C07K14/47 C12Q1/68

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 7 C12Q

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)
SEQUENCE SEARCH, BIOSIS, MEDLINE, EPO-Internal, WPI Data, EMBASE

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	BERTUCCI F ET AL: "EXPRESSION SCANNING OF AN ARRAY OF GROWTH CONTROL GENES IN HUMAN TUMOR CELL LINES" ONCOGENE, BASINGSTOKE, HANTS, GB, vol. 18, no. 26, 1999, pages 3905-3912, XP000979482 ISSN: 0950-9232 the whole document	1-42
X	PEROU C M ET AL: "Molecular portraits of human breast tumours." NATURE. ENGLAND 17 AUG 2000, vol. 406, no. 6797, 17 August 2000 (2000-08-17), pages 747-752, XP002235791 ISSN: 0028-0836 the whole document	1-42



Further documents are listed in the continuation of box C.



Patent family members are listed in annex.

* Special categories of cited documents :

- "A" document defining the general state of the art which is not considered to be of particular relevance
- "E" earlier document but published on or after the international filing date
- "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- "O" document referring to an oral disclosure, use, exhibition or other means
- "P" document published prior to the international filing date but later than the priority date claimed

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"&" document member of the same patent family

Date of the actual completion of the international search

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Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax: (+31-70) 340-3016

Authorized officer

Luzzatto, E

INTERNATIONAL SEARCH REPORT

Int'l Application No

PCI/18 01/02811

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	HOCH R V ET AL: "GATA-3 is expressed in association with estrogen receptor in breast cancer." INTERNATIONAL JOURNAL OF CANCER. JOURNAL INTERNATIONAL DU CANCER. UNITED STATES 20 APR 1999, vol. 84, no. 2, 20 April 1999 (1999-04-20), pages 122-128, XP001146467 ISSN: 0020-7136	1-5,8,9, 12,13, 16,17, 20,21, 24-42
Y	the whole document	6,7,10, 11,14, 15,18, 19,22,23
Y	----- DATABASE MEDLINE 'Online! US NATIONAL LIBRARY OF MEDICINE (NLM), BETHESDA, MD, US; July 1997 (1997-07) SHAW-BRUHA C M ET AL: "Expression of the prolactin gene in normal and neoplastic human breast tissues and human mammary cell lines: promoter usage and alternative mRNA splicing." Database accession no. NLM9266104 XP002235792 abstract & BREAST CANCER RESEARCH AND TREATMENT. NETHERLANDS JUL 1997, vol. 44, no. 3, July 1997 (1997-07), pages 243-253, ISSN: 0167-6806	6,7
Y	----- GRAHAM J D ET AL: "Regulation of the expression and activity by progestins of a member of the SOX gene family of transcriptional modulators." JOURNAL OF MOLECULAR ENDOCRINOLOGY. ENGLAND JUN 1999, vol. 22, no. 3, June 1999 (1999-06), pages 295-304, XP000995364 ISSN: 0952-5041 the whole document	10,11, 14,15, 18,19
Y	----- MAGUIRE T M ET AL: "High levels of cathepsin B predict poor outcome in patients with breast cancer." INTERNATIONAL JOURNAL OF BIOLOGICAL MARKERS, vol. 13, no. 3, July 1998 (1998-07), pages 139-144, XP001118861 ISSN: 0393-6155 abstract	22,23
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INTERNATIONAL SEARCH REPORT

Int. Application No

PCT/IB 01/02811

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	MATHOULIN-PORTIER M P ET AL: "Prognostic value of simultaneous expression of p21 and mdm2 in breast carcinomas treated by adjuvant chemotherapy with anthracyclin." ONCOLOGY REPORTS. GREECE 2000 MAY-JUN, vol. 7, no. 3, May 2000 (2000-05), pages 675-680, XP009007614 ISSN: 1021-335X the whole document ---	1-42
A	FAIRCHILD C R ET AL: "ISOLATION OF AMPLIFIED AND OVEREXPRESSED DNA SEQUENCES FROM ADRIAMYCIN-RESISTANT HUMAN BREAST CANCER CELLS" CANCER RESEARCH, vol. 47, no. 19, 1987, pages 5141-5148, XP009006931 ISSN: 0008-5472 the whole document ---	1-42
A	VARGAS-ROIG L M ET AL: "c-erbB-2 (HER-2/neu) protein and drug resistance in breast cancer patients treated with induction chemotherapy." INTERNATIONAL JOURNAL OF CANCER. JOURNAL INTERNATIONAL DU CANCER. UNITED STATES 20 APR 1999, vol. 84, no. 2, 20 April 1999 (1999-04-20), pages 129-134, XP002235684 ISSN: 0020-7136 the whole document ---	1-42
A	PENAULT-LLORCA F ET AL: "Expression of FGF and FGF receptor genes in human breast cancer." INTERNATIONAL JOURNAL OF CANCER. JOURNAL INTERNATIONAL DU CANCER. UNITED STATES 10 APR 1995, vol. 61, no. 2, 10 April 1995 (1995-04-10), pages 170-176, XP009007615 ISSN: 0020-7136 the whole document ---	1-42
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INTERNATIONAL SEARCH REPORT

Int. Application No

PCT/IB 01/02811

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	<p>DATABASE EBI 'Online! Homo sapiens cathepsin B mRNA, complete cds., 20 May 1993 (1993-05-20) CAO L. ET AL.: Database accession no. L16510 XP002235958 abstract & CAO L. ET AL.: "Human gastric adenocarcinoma cathepsin B: isolation and sequencing of full-length cDNAs and polymorphism of the gene" GENE, vol. 139, no. 2, 1994, pages 163-169,</p>	22,23
A	<p>CAFFO O ET AL: "Prognostic value of p21(WAF1) and p53 expression in breast carcinoma: an immunohistochemical study in 261 patients with long-term follow-up." CLINICAL CANCER RESEARCH: AN OFFICIAL JOURNAL OF THE AMERICAN ASSOCIATION FOR CANCER RESEARCH. UNITED STATES SEP 1996, vol. 2, no. 9, September 1996 (1996-09), pages 1591-1599, XP001121315 ISSN: 1078-0432 the whole document</p>	1-42
T	<p>BERTUCCI FRANÇOIS ET AL: "Gene expression profiles of poor-prognosis primary breast cancer correlate with survival." HUMAN MOLECULAR GENETICS. ENGLAND 15 APR 2002, vol. 11, no. 8, 15 April 2002 (2002-04-15), pages 863-872, XP001146244 ISSN: 0964-6906</p>	

INTERNATIONAL SEARCH REPORT

ational application No.
PCT/IB 01/02811**Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)**

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☒ Claims Nos.: 6,7,10,11,14,15,18,19,22-26,28-43 (all partly)
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
see FURTHER INFORMATION sheet PCT/ISA/210
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

see additional sheet

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
1-43 all partially

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 1-43 (all partially)

Polynucleotide libraries comprising any of SEQ IDs 76-78, which according to table 10 of the application correspond to the GATA-3 gene, method for detecting cancer correlated differentially expressed polynucleotide sequences or for screening an anti-tumour agent, comprising the step of reacting a polynucleotide sample from a patient with a probe comprising any of SEQ IDs 76-78.

2. Claims: 1-3, 8-11, 16-27,29,31-43 (all partially)

Polynucleotide libraries comprising any of SEQ IDs 354-355, which according to table 10 of the application correspond to the MYB gene, method for detecting cancer correlated differentially expressed polynucleotide sequences or for screening an anti-tumour agent, comprising the step of reacting a polynucleotide sample from a patient with a probe comprising any of SEQ IDs 354-355.

3. Claims: 1-3, 8-13,27,29,30,33-43 (partially)

Polynucleotide libraries comprising any of SEQ IDs 322-323, which according to table 10 of the application correspond to the KIAA1075 gene, method for detecting cancer correlated differentially expressed polynucleotide sequences or for screening an anti-tumour agent, comprising the step of reacting a polynucleotide sample from a patient with a probe comprising any of SEQ IDs 322-323.

Inventions 4-171: subject as in inventions 1-3, however the SEQ IDs are those corresponding to genes 4-171 as listed in table 10.

As to the claims related to each of inventions 4-171, the ISA wishes to draw the applicant's attention to the following:

In view of the number of genes and sequences to which the claims relate, and of the fact that the vast majority of SEQ IDs are mentioned only in some claims and not in others, it becomes unduly burdensome for the ISA to provide for each invention the indication as to which claims are related to it. Should any additional search fees be paid, the clear definition of each invention by means of the SEQ IDs will allow the unambiguous definition of the claims that belong to the invention to be searched.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Continuation of Box I.2

Claims Nos.: 6,7,10,11,14,15,18,19,22-26,28-43 (all partly)

The present remarks apply to the only invention searched, i.e. invention 1 (see "Invitation to pay additional fees"). Should further search fees be paid, similar remarks could apply to the further inventions to be searched, leading to an incomplete search also with respect to these inventions.

1) The application relates to genes which are differentially expressed in breast tumour cells compared to normal breast cells (see p. 21, l. 15-p. 22, l. 12). No other cells are analysed for differential gene expression of tumor vs. normal cells. However, claims 33-40 and 42-43 encompass the detection of differentially expressed sequences correlated with "a cancer", i.e. they are not limited to breast cancer. Hence, they lack support (Art. 6 PCT) to such an extent as to render a meaningful search over their whole scope impossible. Claims 33-40 and 42-43 have therefore been searched only insofar as related to breast cancer.

2) Claims 6,10,14,18,22 relate to a very large number of combinations comprising the GATA-3 gene related sequences and at least one other of the second group of sequences to which the said claims relate. They therefore lack conciseness (Art. 6 PCT) to such an extent as to render impossible a meaningful search covering all the said combinations. The search has thus been limited to combinations comprising the GATA-3 related sequences of set 32 (consisting of SEQ IDs 76-78) and those of sets 38, 11 and 14 (SEQ IDs 91-93, 22-24 and 30-31 respectively), these being the first sets mentioned in the various claims in the context of combinations with the GATA-3 gene. The same applies to claims 7,11,15,19,23-26,28-43 insofar as dependent on the said claims.

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.